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<110> Schupp, Thomas
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Molnar, Istvan
Zirkle, Ross
Cyr, Devon
Goerlach, Joern

<120> GENES FOR THE BIOSYNTHESIS OF EPOTHILONES

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 1045 1050 1055
 Arg Pro Gln Ala Leu Ser Ala Ile Glu Ala Ala Ala Met Pro Val Ala
 1060 1065 1070
 Tyr Leu Thr Ala Trp Tyr Ala Leu Asp Arg Ile Ala Arg Leu Gln Pro
 1075 1080 1085
 Gly Glu Arg Val Leu Ile His Ala Ala Thr Gly Gly Val Gly Leu Ala
 1090 1095 1100

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Ala Val Gln Trp Ala Gln His Val Gly Ala Glu Val His Ala Thr Ala
 1105 1110 1115 1120

Gly Thr Pro Glu Lys Arg Ala Tyr Leu Glu Ser Leu Gly Val Arg Tyr
 1125 1130 1135

Val Ser Asp Ser Arg Ser Asp Arg Phe Val Ala Asp Val Arg Ala Trp
 1140 1145 1150

Thr Gly Gly Glu Gly Val Asp Val Val Leu Asn Ser Leu Ser Gly Glu
 1155 1160 1165

Leu Ile Asp Lys Ser Phe Asn Leu Leu Arg Ser His Gly Arg Phe Val
 1170 1175 1180

Glu Leu Gly Lys Arg Asp Cys Tyr Ala Asp Asn Gln Leu Gly Leu Arg
 1185 1190 1195 1200

Pro Phe Leu Arg Asn Leu Ser Phe Ser Leu Val Asp Leu Arg Gly Met
 1205 1210 1215

Met Leu Glu Arg Pro Ala Arg Val Arg Ala Leu Leu Glu Glu Leu Leu
 1220 1225 1230

Gly Leu Ile Ala Ala Gly Val Phe Thr Pro Pro Pro Ile Ala Thr Leu
 1235 1240 1245

Pro Ile Ala Arg Val Ala Asp Ala Phe Arg Ser Met Ala Gln Ala Gln
 1250 1255 1260

His Leu Gly Lys Leu Val Leu Thr Leu Gly Asp Pro Glu Val Gln Ile
 1265 1270 1275 1280

Arg Ile Pro Thr His Ala Gly Ala Gly Pro Ser Thr Gly Asp Arg Asp
 1285 1290 1295

Leu Leu Asp Arg Leu Ala Ser Ala Ala Pro Ala Ala Arg Ala Ala Ala
 1300 1305 1310

Leu Glu Ala Phe Leu Arg Thr Gln Val Ser Gln Val Leu Arg Thr Pro
 1315 1320 1325

Glu Ile Lys Val Gly Ala Glu Ala Leu Phe Thr Arg Leu Gly Met Asp
 1330 1335 1340

Ser Leu Met Ala Val Glu Leu Arg Asn Arg Ile Glu Ala Ser Leu Lys
 1345 1350 1355 1360

Leu Lys Leu Ser Thr Thr Phe Leu Ser Thr Ser Pro Asn Ile Ala Leu
 1365 1370 1375

Leu Ala Gln Asn Leu Leu Asp Ala Leu Ala Thr Ala Leu Ser Leu Glu
 1380 1385 1390

Arg Val Ala Ala Glu Asn Leu Arg Ala Gly Val Gln Asn Asp Phe Val
 1395 1400 1405

Ser Ser Gly Ala Asp Gln Asp Trp Glu Ile Ile Ala Leu
 1410 1415 1420

<210> 3

<211> 1410

<212> PRT

<213> Sorangium cellulosum

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<400> 3

Met Thr Ile Asn Gln Leu Leu Asn Glu Leu Glu His Gln Gly Ile Lys
 1 5 10 15
 Leu Ala Ala Asp Gly Glu Arg Leu Gln Ile Gln Ala Pro Lys Asn Ala
 20 25 30
 Leu Asn Pro Asn Leu Leu Ala Arg Ile Ser Glu His Lys Ser Thr Ile
 35 40 45
 Leu Thr Met Leu Arg Gln Arg Leu Pro Ala Glu Ser Ile Val Pro Ala
 50 55 60
 Pro Ala Glu Arg His Ala Pro Phe Pro Leu Thr Asp Ile Gln Glu Ser
 65 70 75 80
 Tyr Trp Leu Gly Arg Thr Gly Ala Phe Thr Val Pro Ser Gly Ile His
 85 90 95
 Ala Tyr Arg Glu Tyr Asp Cys Thr Asp Leu Asp Val Pro Arg Leu Ser
 100 105 110
 Arg Ala Phe Arg Lys Val Val Ala Arg His Asp Met Leu Arg Ala His
 115 120 125
 Thr Leu Pro Asp Met Met Gln Val Ile Glu Pro Lys Val Asp Ala Asp
 130 135 140
 Ile Glu Ile Ile Asp Leu Arg Gly Leu Asp Arg Ser Thr Arg Glu Ala
 145 150 155 160
 Arg Leu Val Ser Leu Arg Asp Ala Met Ser His Arg Ile Tyr Asp Thr
 165 170 175
 Glu Arg Pro Pro Leu Tyr His Val Val Ala Val Arg Leu Asp Glu Arg
 180 185 190
 Gln Thr Arg Leu Val Leu Ser Ile Asp Leu Ile Asn Val Asp Leu Gly
 195 200 205
 Ser Leu Ser Ile Ile Phe Lys Asp Trp Leu Ser Phe Tyr Glu Asp Pro
 210 215 220
 Glu Thr Ser Leu Pro Val Leu Glu Leu Ser Tyr Arg Asp Tyr Val Leu
 225 230 235 240
 Ala Leu Glu Ser Arg Lys Lys Ser Glu Ala His Gln Arg Ser Met Asp
 245 250 255
 Tyr Trp Lys Arg Arg Ile Ala Glu Leu Pro Pro Pro Pro Thr Leu Pro
 260 265 270
 Met Lys Ala Asp Pro Ser Thr Leu Lys Glu Ile Arg Phe Arg His Thr
 275 280 285
 Glu Gln Trp Leu Pro Ser Asp Ser Trp Gly Arg Leu Lys Arg Arg Val
 290 295 300
 Gly Glu Arg Gly Leu Thr Pro Thr Gly Val Ile Leu Ala Ala Phe Ser
 305 310 315 320
 Glu Val Ile Gly Arg Trp Ser Ala Ser Pro Arg Phe Thr Leu Asn Ile
 325 330 335
 Thr Leu Phe Asn Arg Leu Pro Val His Pro Arg Val Asn Asp Ile Thr

340	345	350
Gly Asp Phe Thr Ser Met Val	Leu Leu Asp Ile Asp Thr Thr Arg Asp	
355	360	365
Lys Ser Phe Glu Gln Arg Ala Lys Arg Ile Gln Glu Gln Leu Trp Glu		
370	375	380
Ala Met Asp His Cys Asp Val Ser Gly Ile Glu Val Gln Arg Glu Ala		
385	390	395
Ala Arg Val Leu Gly Ile Gln Arg Gly Ala Leu Phe Pro Val Val Leu		
405	410	415
Thr Ser Ala Leu Asn Gln Gln Val Val Gly Val Thr Ser Leu Gln Arg		
420	425	430
Leu Gly Thr Pro Val Tyr Thr Ser Thr Gln Thr Pro Gln Leu Leu Leu		
435	440	445
Asp His Gln Leu Tyr Glu His Asp Gly Asp Leu Val Leu Ala Trp Asp		
450	455	460
Ile Val Asp Gly Val Phe Pro Pro Asp Leu Leu Asp Asp Met Leu Glu		
465	470	475
Ala Tyr Val Val Phe Leu Arg Arg Leu Thr Glu Glu Pro Trp Gly Glu		
485	490	495
Gln Val Arg Cys Ser Leu Pro Pro Ala Gln Leu Glu Ala Arg Ala Ser		
500	505	510
Ala Asn Ala Thr Asn Ala Leu Leu Ser Glu His Thr Leu His Gly Leu		
515	520	525
Phe Ala Ala Arg Val Glu Gln Leu Pro Met Gln Leu Ala Val Val Ser		
530	535	540
Ala Arg Lys Thr Leu Thr Tyr Glu Glu Leu Ser Arg Arg Ser Arg Arg		
545	550	555
Leu Gly Ala Arg Leu Arg Glu Gln Gly Ala Arg Pro Asn Thr Leu Val		
565	570	575
Ala Val Val Met Glu Lys Gly Trp Glu Gln Val Val Ala Val Leu Ala		
580	585	590
Val Leu Glu Ser Gly Ala Ala Tyr Val Pro Ile Asp Ala Asp Leu Pro		
595	600	605
Ala Glu Arg Ile His Tyr Leu Leu Asp His Gly Glu Val Lys Leu Val		
610	615	620
Leu Thr Gln Pro Trp Leu Asp Gly Lys Leu Ser Trp Pro Pro Gly Ile		
625	630	635
Gln Arg Leu Leu Val Ser Glu Ala Gly Val Glu Gly Asp Gly Asp Gln		
645	650	655
Pro Pro Met Met Pro Ile Gln Thr Pro Ser Asp Leu Ala Tyr Val Ile		
660	665	670
Tyr Thr Ser Gly Ser Thr Gly Leu Pro Lys Gly Val Met Ile Asp His		
675	680	685
Arg Gly Ala Val Asn Thr Ile Leu Asp Ile Asn Glu Arg Phe Glu Ile		

690	695	700
Gly 705	Pro Gly Asp Arg Val 710	Leu Ala Leu Ser Ser 715
Ser 720	Val Phe Gly Ile Leu 725	Ala Ala Gly Gly Thr 730
Val 735	Pro Asp Ala Ser Lys Leu Arg 740	Pro Ala His Trp Ala 745
Ile 750	Glu Arg Glu Lys Val Thr 755	Val Trp Asn Ser Val 760
Arg 765	Met Leu Val Glu His Phe 770	Glu Gly Arg Pro Asp 775
Ser 780	Leu Arg Leu Ser Leu 785	Leu Ser Gly Asp Trp 790
Pro 800	Gly Glu Leu Gln Ala Ile 805	Val Ser Val Ile Ser 810
Gly 815	Gly Ala Thr Glu Ala Ser 820	Ile Trp Ser Ile Gly 825
Asn 830	Val Asp Leu Ser Trp Ala 835	Ile Pro Tyr Gly Arg 840
Asn 845	Gln Thr Phe His Val 850	Leu Asp Glu Ala Leu 855
Trp 860	Val Pro Gly Gln Leu Tyr 865	Ile Gly Gly Val Gly 870
Tyr 875	Trp Arg Asp Glu Glu Lys 880	Thr Arg Lys Ser Phe 885
Glu 890	Thr Gly Glu Arg Leu Tyr 900	Lys Thr Gly Asp Leu 905
Pro 910	Asp Gly Asn Ile Glu Phe 915	Arg Glu Asp Asn Gln 920
Leu 925	Arg Gly Tyr Arg Val Glu 930	Leu Gly Glu Ile Glu 935
Ser 940	His Pro Asn Val Arg Asp 945	Ala Val Ile Val Pro 950
Ala 955	Ala Asn Lys Leu Leu Leu 960	Ala Tyr Val Val Pro 965
Arg 970	Arg Ala Ala Glu Gln Asp 975	Ala Ser Leu Lys Thr 980
Ala 985	Arg Ala His Ala Ala Glu 990	Ala Asp Gly Leu Ser 995
Val 1000	Gln Phe Lys Leu Ala Arg 1005	His Gly Leu Arg Arg 1010
Lys 1015	Pro Val Val Asp Leu Thr 1020	Gly Gln Asp Pro Arg 1025
Asp 1030	Val Tyr Ala Arg Arg Arg 1035	Ser Val Arg Thr Phe 1040
		Leu Glu Ala Pro

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1045	1050	1055
Ile Pro Phe Val Glu Phe Gly Arg Phe Leu Ser Cys Leu Ser Ser Val 1060	1065	1070
Glu Pro Asp Gly Ala Thr Leu Pro Lys Phe Arg Tyr Pro Ser Ala Gly 1075	1080	1085
Ser Thr Tyr Pro Val Gln Thr Tyr Ala Tyr Val Lys Ser Gly Arg Ile 1090	1095	1100
Glu Gly Val Asp Glu Gly Phe Tyr Tyr Tyr His Pro Phe Glu His Arg 1105	1110	1115
Leu Leu Lys Leu Ser Asp His Gly Ile Glu Arg Gly Ala His Val Arg 1125	1130	1135
Gln Asn Phe Asp Val Phe Asp Glu Ala Ala Phe Asn Leu Leu Phe Val 1140	1145	1150
Gly Arg Ile Asp Ala Ile Glu Ser Leu Tyr Gly Ser Ser Ser Arg Glu 1155	1160	1165
Phe Cys Leu Leu Glu Ala Gly Tyr Met Ala Gln Leu Leu Met Glu Gln 1170	1175	1180
Ala Pro Ser Cys Asn Ile Gly Val Cys Pro Val Gly Gln Phe Asn Phe 1185	1190	1195
Glu Gln Val Arg Pro Val Leu Asp Leu Arg His Ser Asp Val Tyr Val 1205	1210	1215
His Gly Met Leu Gly Gly Arg Val Asp Pro Arg Gln Phe Gln Val Cys 1220	1225	1230
Thr Leu Gly Gln Asp Ser Ser Pro Arg Arg Ala Thr Thr Arg Gly Ala 1235	1240	1245
Pro Pro Gly Arg Glu Gln His Phe Ala Asp Met Leu Arg Asp Phe Leu 1250	1255	1260
Arg Thr Lys Leu Pro Glu Tyr Met Val Pro Thr Val Phe Val Glu Leu 1265	1270	1275
Asp Ala Leu Pro Leu Thr Ser Asn Gly Lys Val Asp Arg Lys Ala Leu 1285	1290	1295
Arg Glu Arg Lys Asp Thr Ser Ser Pro Arg His Ser Gly His Thr Ala 1300	1305	1310
Pro Arg Asp Ala Leu Glu Glu Ile Leu Val Ala Val Val Arg Glu Val 1315	1320	1325
Leu Gly Leu Glu Val Val Gly Leu Gln Gln Ser Phe Val Asp Leu Gly 1330	1335	1340
Ala Thr Ser Ile His Ile Val Arg Met Arg Ser Leu Leu Gln Lys Arg 1345	1350	1355
Leu Asp Arg Glu Ile Ala Ile Thr Glu Leu Phe Gln Tyr Pro Asn Leu 1365	1370	1375
Gly Ser Leu Ala Ser Gly Leu Arg Arg Asp Ser Arg Asp Leu Asp Gln 1380	1385	1390
Arg Pro Asn Met Gln Asp Arg Val Glu Val Arg Arg Lys Gly Arg Arg		

1395

1400

1405

Arg Ser
1410

<210> 4

<211> 1832

<212> PRT

<213> Sorangium cellulosum

<400> 4

Met Glu Glu Gln Glu Ser Ser Ala Ile Ala Val Ile Gly Met Ser Gly
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20 25 30Asp Gly Thr Glu Ala Val Gln Arg Phe Ser Glu Gln Glu Leu Ala Ala
35 40 45Ser Gly Val Asp Pro Ala Leu Val Leu Asp Pro Ser Tyr Val Arg Ala
50 55 60Gly Ser Val Leu Glu Asp Val Asp Arg Phe Asp Ala Ala Phe Phe Gly
65 70 75 80Ile Ser Pro Arg Glu Ala Glu Leu Met Asp Pro Gln His Arg Ile Phe
85 90 95Met Glu Cys Ala Trp Glu Ala Leu Glu Asn Ala Gly Tyr Asp Pro Thr
100 105 110Ala Tyr Glu Gly Ser Ile Gly Val Tyr Ala Gly Ala Asn Met Ser Ser
115 120 125Tyr Leu Thr Ser Asn Leu His Glu His Pro Ala Met Met Arg Trp Pro
130 135 140Gly Trp Phe Gln Thr Leu Ile Gly Asn Asp Lys Asp Tyr Leu Ala Thr
145 150 155 160His Val Ser Tyr Arg Leu Asn Leu Arg Gly Pro Ser Ile Ser Val Gln
165 170 175Thr Ala Cys Ser Thr Ser Leu Val Ala Val His Leu Ala Cys Met Ser
180 185 190Leu Leu Asp Arg Glu Cys Asp Met Ala Leu Ala Gly Gly Ile Thr Val
195 200 205Arg Ile Pro His Arg Ala Gly Tyr Val Tyr Ala Glu Gly Gly Ile Phe
210 215 220Ser Pro Asp Gly His Cys Arg Ala Phe Asp Ala Lys Ala Asn Gly Thr
225 230 235 240Ile Met Gly Asn Gly Cys Gly Val Val Leu Leu Lys Pro Leu Asp Arg
245 250 255Ala Leu Ser Asp Gly Asp Pro Val Arg Ala Val Ile Leu Gly Ser Ala
260 265 270Thr Asn Asn Asp Gly Ala Arg Lys Ile Gly Phe Thr Ala Pro Ser Glu
275 280 285

Val Gly Gln Ala Gln Ala Ile Met Glu Ala Leu Ala Leu Ala Gly Val
 290 295 300
 Glu Ala Arg Ser Ile Gln Tyr Ile Glu Thr His Gly Thr Gly Thr Leu
 305 310 315 320
 Leu Gly Asp Ala Ile Glu Thr Ala Ala Leu Arg Arg Val Phe Gly Arg
 325 330 335
 Asp Ala Ser Ala Arg Arg Ser Cys Ala Ile Gly Ser Val Lys Thr Gly
 340 345 350
 Ile Gly His Leu Glu Ser Ala Ala Gly Ile Ala Gly Leu Ile Lys Thr
 355 360 365
 Val Leu Ala Leu Glu His Arg Gln Leu Pro Pro Ser Leu Asn Phe Glu
 370 375 380
 Ser Pro Asn Pro Ser Ile Asp Phe Ala Ser Ser Pro Phe Tyr Val Asn
 385 390 395 400
 Thr Ser Leu Lys Asp Trp Asn Thr Gly Ser Thr Pro Arg Arg Ala Gly
 405 410 415
 Val Ser Ser Phe Gly Ile Gly Gly Thr Asn Ala His Val Val Leu Glu
 420 425 430
 Glu Ala Pro Ala Ala Lys Leu Pro Ala Ala Ala Pro Ala Arg Ser Ala
 435 440 445
 Glu Leu Phe Val Val Ser Ala Lys Ser Ala Ala Ala Leu Asp Ala Ala
 450 455 460
 Ala Ala Arg Leu Arg Asp His Leu Gln Ala His Gln Gly Ile Ser Leu
 465 470 475 480
 Gly Asp Val Ala Phe Ser Leu Ala Thr Thr Arg Ser Pro Met Glu His
 485 490 495
 Arg Leu Ala Met Ala Ala Pro Ser Arg Glu Ala Leu Arg Glu Gly Leu
 500 505 510
 Asp Ala Ala Ala Arg Gly Gln Thr Pro Pro Gly Ala Val Arg Gly Arg
 515 520 525
 Cys Ser Pro Gly Asn Val Pro Lys Val Val Phe Val Phe Pro Gly Gln
 530 535 540
 Gly Ser Gln Trp Val Gly Met Gly Arg Gln Leu Leu Ala Glu Glu Pro
 545 550 555 560
 Val Phe His Ala Ala Leu Ser Ala Cys Asp Arg Ala Ile Gln Ala Glu
 565 570 575
 Ala Gly Trp Ser Leu Leu Ala Glu Leu Ala Ala Asp Glu Gly Ser Ser
 580 585 590
 Gln Leu Glu Arg Ile Asp Val Val Gln Pro Val Leu Phe Ala Leu Ala
 595 600 605
 Val Ala Phe Ala Ala Leu Trp Arg Ser Trp Gly Val Ala Pro Asp Val
 610 615 620
 Val Ile Gly His Ser Met Gly Glu Val Ala Ala Ala His Val Ala Gly
 625 630 635 640

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Ala Leu Ser Leu Glu Asp Ala Val Ala Ile Ile Cys Arg Arg Ser Arg
 645 650 655
 Leu Leu Arg Arg Ile Ser Gly Gln Gly Glu Met Ala Val Thr Glu Leu
 660 665 670
 Ser Leu Ala Glu Ala Glu Ala Ala Leu Arg Gly Tyr Glu Asp Arg Val
 675 680 685
 Ser Val Ala Val Ser Asn Ser Pro Arg Ser Thr Val Leu Ser Gly Glu
 690 695 700
 Pro Ala Ala Ile Gly Glu Val Leu Ser Ser Leu Asn Ala Lys Gly Val
 705 710 715 720
 Phe Cys Arg Arg Val Lys Val Asp Val Ala Ser His Ser Pro Gln Val
 725 730 735
 Asp Pro Leu Arg Glu Asp Leu Leu Ala Ala Leu Gly Gly Leu Arg Pro
 740 745 750
 Gly Ala Ala Ala Val Pro Met Arg Ser Thr Val Thr Gly Ala Met Val
 755 760 765
 Ala Gly Pro Glu Leu Gly Ala Asn Tyr Trp Met Asn Asn Leu Arg Gln
 770 775 780
 Pro Val Arg Phe Ala Glu Val Val Gln Ala Gln Leu Gln Gly Gly His
 785 790 795 800
 Gly Leu Phe Val Glu Met Ser Pro His Pro Ile Leu Thr Thr Ser Val
 805 810 815
 Glu Glu Met Arg Arg Ala Ala Gln Arg Ala Gly Ala Ala Val Gly Ser
 820 825 830
 Leu Arg Arg Gly Gln Asp Glu Arg Pro Ala Met Leu Glu Ala Leu Gly
 835 840 845
 Thr Leu Trp Ala Gln Gly Tyr Pro Val Pro Trp Gly Arg Leu Phe Pro
 850 855 860
 Ala Gly Gly Arg Arg Val Pro Leu Pro Thr Tyr Pro Trp Gln Arg Glu
 865 870 875 880
 Arg Tyr Trp Ile Glu Ala Pro Ala Lys Ser Ala Ala Gly Asp Arg Arg
 885 890 895
 Gly Val Arg Ala Gly Gly His Pro Leu Leu Gly Glu Met Gln Thr Leu
 900 905 910
 Ser Thr Gln Thr Ser Thr Arg Leu Trp Glu Thr Thr Leu Asp Leu Lys
 915 920 925
 Arg Leu Pro Trp Leu Gly Asp His Arg Val Gln Gly Ala Val Val Phe
 930 935 940
 Pro Gly Ala Ala Tyr Leu Glu Met Ala Ile Ser Ser Gly Ala Glu Ala
 945 950 955 960
 Leu Gly Asp Gly Pro Leu Gln Ile Thr Asp Val Val Leu Ala Glu Ala
 965 970 975
 Leu Ala Phe Ala Gly Asp Ala Ala Val Leu Val Gln Val Val Thr Thr
 980 985 990

Glu Gln Pro Ser Gly Arg Leu Gln Phe Gln Ile Ala Ser Arg Ala Pro
 995 1000 1005
 Gly Ala Gly His Ala Ser Phe Arg Val His Ala Arg Gly Ala Leu Leu
 1010 1015 1020
 Arg Val Glu Arg Thr Glu Val Pro Ala Gly Leu Thr Leu Ser Ala Val
 1025 1030 1035 1040
 Arg Ala Arg Leu Gln Ala Ser Ile Pro Ala Ala Ala Thr Tyr Ala Glu
 1045 1050 1055
 Leu Thr Glu Met Gly Leu Gln Tyr Gly Pro Ala Phe Gln Gly Ile Ala
 1060 1065 1070
 Glu Leu Trp Arg Gly Glu Gly Glu Ala Leu Gly Arg Val Arg Leu Pro
 1075 1080 1085
 Asp Ala Ala Gly Ser Ala Ala Glu Tyr Arg Leu His Pro Ala Leu Leu
 1090 1095 1100
 Asp Ala Cys Phe Gln Ile Val Gly Ser Leu Phe Ala Arg Ser Gly Glu
 1105 1110 1115 1120
 Ala Thr Pro Trp Val Pro Val Glu Leu Gly Ser Leu Arg Leu Leu Gln
 1125 1130 1135
 Arg Pro Ser Gly Glu Leu Trp Cys His Ala Arg Val Val Asn His Gly
 1140 1145 1150
 His Gln Thr Pro Asp Arg Gln Gly Ala Asp Phe Trp Val Val Asp Ser
 1155 1160 1165
 Ser Gly Ala Val Val Ala Glu Val Cys Gly Leu Val Ala Gln Arg Leu
 1170 1175 1180
 Pro Gly Gly Val Arg Arg Arg Glu Glu Asp Asp Trp Phe Leu Glu Leu
 1185 1190 1195 1200
 Glu Trp Glu Pro Ala Ala Val Gly Thr Ala Lys Val Asn Ala Gly Arg
 1205 1210 1215
 Trp Leu Leu Leu Gly Gly Gly Gly Gly Leu Gly Ala Ala Leu Arg Ala
 1220 1225 1230
 Met Leu Glu Ala Gly Gly His Ala Val Val His Ala Ala Glu Asn Asn
 1235 1240 1245
 Thr Ser Ala Ala Gly Val Arg Ala Leu Leu Ala Lys Ala Phe Asp Gly
 1250 1255 1260
 Gln Ala Pro Thr Ala Val Val His Leu Gly Ser Leu Asp Gly Gly Gly
 1265 1270 1275 1280
 Glu Leu Asp Pro Gly Leu Gly Ala Gln Gly Ala Leu Asp Ala Pro Arg
 1285 1290 1295
 Ser Ala Asp Val Ser Pro Asp Ala Leu Asp Pro Ala Leu Val Arg Gly
 1300 1305 1310
 Cys Asp Ser Val Leu Trp Thr Val Gln Ala Leu Ala Gly Met Gly Phe
 1315 1320 1325
 Arg Asp Ala Pro Arg Leu Trp Leu Leu Thr Arg Gly Ala Gln Ala Val
 1330 1335 1340

Gly Ala Gly Asp Val Ser Val Thr Gln Ala Pro Leu Leu Gly Leu Gly
 1345 1350 1355 1360
 Arg Val Ile Ala Met Glu His Ala Asp Leu Arg Cys Ala Arg Val Asp
 1365 1370 1375
 Leu Asp Pro Ala Arg Pro Glu Gly Glu Leu Ala Ala Leu Leu Ala Glu
 1380 1385 1390
 Leu Leu Ala Asp Asp Ala Glu Ala Glu Val Ala Leu Arg Gly Gly Glu
 1395 1400 1405
 Arg Cys Val Ala Arg Ile Val Arg Arg Gln Pro Glu Thr Arg Pro Arg
 1410 1415 1420
 Gly Arg Ile Glu Ser Cys Val Pro Thr Asp Val Thr Ile Arg Ala Asp
 1425 1430 1435 1440
 Ser Thr Tyr Leu Val Thr Gly Gly Leu Gly Gly Leu Gly Leu Ser Val
 1445 1450 1455
 Ala Gly Trp Leu Ala Glu Arg Gly Ala Gly His Leu Val Leu Val Gly
 1460 1465 1470
 Arg Ser Gly Ala Ala Ser Val Glu Gln Arg Ala Ala Val Ala Ala Leu
 1475 1480 1485
 Glu Ala Arg Gly Ala Arg Val Thr Val Ala Lys Ala Asp Val Ala Asp
 1490 1495 1500
 Arg Ala Gln Leu Glu Arg Ile Leu Arg Glu Val Thr Thr Ser Gly Met
 1505 1510 1515 1520
 Pro Leu Arg Gly Val Val His Ala Ala Gly Ile Leu Asp Asp Gly Leu
 1525 1530 1535
 Leu Met Gln Gln Thr Pro Ala Arg Phe Arg Lys Val Met Ala Pro Lys
 1540 1545 1550
 Val Gln Gly Ala Leu His Leu His Ala Leu Thr Arg Glu Ala Pro Leu
 1555 1560 1565
 Ser Phe Phe Val Leu Tyr Ala Ser Gly Val Gly Leu Leu Gly Ser Pro
 1570 1575 1580
 Gly Gln Gly Asn Tyr Ala Ala Ala Asn Thr Phe Leu Asp Ala Leu Ala
 1585 1590 1595 1600
 His His Arg Arg Ala Gln Gly Leu Pro Ala Leu Ser Val Asp Trp Gly
 1605 1610 1615
 Leu Phe Ala Glu Val Gly Met Ala Ala Ala Gln Glu Asp Arg Gly Ala
 1620 1625 1630
 Arg Leu Val Ser Arg Gly Met Arg Ser Leu Thr Pro Asp Glu Gly Leu
 1635 1640 1645
 Ser Ala Leu Ala Arg Leu Leu Glu Ser Gly Arg Ala Gln Val Gly Val
 1650 1655 1660
 Met Pro Val Asn Pro Arg Leu Trp Val Glu Leu Tyr Pro Ala Ala Ala
 1665 1670 1675 1680
 Ser Ser Arg Met Leu Ser Arg Leu Val Thr Ala His Arg Ala Ser Ala
 1685 1690 1695

Gly Gly Pro Ala Gly Asp Gly Asp Leu Arg Arg Leu Ala Ala Ala
1700 1705 1710

Glu Pro Ser Ala Arg Ser Ala Leu Leu Glu Pro Leu Leu Arg Ala Gln
1715 1720 1725

Ile Ser Gln Val Leu Arg Leu Pro Glu Gly Lys Ile Glu Val Asp Ala
1730 1735 1740

Pro Leu Thr Ser Leu Gly Met Asn Ser Leu Met Gly Leu Glu Leu Arg
1745 1750 1755 1760

Asn Arg Ile Glu Ala Met Leu Gly Ile Thr Val Pro Ala Thr Leu Leu
1765 1770 1775

Trp Thr Tyr Pro Thr Val Ala Ala Leu Ser Gly His Leu Ala Arg Glu
1780 1785 1790

Ala Cys Glu Ala Ala Pro Val Glu Ser Pro His Thr Thr Ala Asp Ser
1795 1800 1805

Ala Val Glu Ile Glu Glu Met Ser Gln Asp Asp Leu Thr Gln Leu Ile
1810 1815 1820

Ala Ala Lys Phe Lys Ala Leu Thr
1825 1830

<210> 5

<211> 7257

<212> PRT

<213> Sorangium cellulosum

<400> 5

Met Thr Thr Arg Gly Pro Thr Ala Gln Gln Asn Pro Leu Lys Gln Ala
1 5 10 15

Ala Ile Ile Ile Gln Arg Leu Glu Glu Arg Leu Ala Gly Leu Ala Gln
20 25 30

Ala Glu Leu Glu Arg Thr Glu Pro Ile Ala Ile Val Gly Ile Gly Cys
35 40 45

Arg Phe Pro Gly Gly Ala Asp Ala Pro Glu Ala Phe Trp Glu Leu Leu
50 55 60

Asp Ala Glu Arg Asp Ala Val Gln Pro Leu Asp Met Arg Trp Ala Leu
65 70 75 80

Val Gly Val Ala Pro Val Glu Ala Val Pro His Trp Ala Gly Leu Leu
85 90 95

Thr Glu Pro Ile Asp Cys Phe Asp Ala Ala Phe Phe Gly Ile Ser Pro
100 105 110

Arg Glu Ala Arg Ser Leu Asp Pro Gln His Arg Leu Leu Leu Glu Val
115 120 125

Ala Trp Glu Gly Leu Glu Asp Ala Gly Ile Pro Pro Arg Ser Ile Asp
130 135 140

Gly Ser Arg Thr Gly Val Phe Val Gly Ala Phe Thr Ala Asp Tyr Ala
145 150 155 160

Arg Thr Val Ala Arg Leu Pro Arg Glu Glu Arg Asp Ala Tyr Ser Ala
165 170 175

Thr Gly Asn Met Leu Ser Ile Ala Ala Gly Arg Leu Ser Tyr Thr Leu
 180 185 190
 Gly Leu Gln Gly Pro Cys Leu Thr Val Asp Thr Ala Cys Ser Ser Ser
 195 200 205
 Leu Val Ala Ile His Leu Ala Cys Arg Ser Leu Arg Ala Gly Glu Ser
 210 215 220
 Asp Leu Ala Leu Ala Gly Gly Val Ser Ala Leu Leu Ser Pro Asp Met
 225 230 235 240
 Met Glu Ala Ala Ala Arg Thr Gln Ala Leu Ser Pro Asp Gly Arg Cys
 245 250 255
 Arg Thr Phe Asp Ala Ser Ala Asn Gly Phe Val Arg Gly Glu Gly Cys
 260 265 270
 Gly Leu Val Val Leu Lys Arg Leu Ser Asp Ala Gln Arg Asp Gly Asp
 275 280 285
 Arg Ile Trp Ala Leu Ile Arg Gly Ser Ala Ile Asn His Asp Gly Arg
 290 295 300
 Ser Thr Gly Leu Thr Ala Pro Asn Val Leu Ala Gln Glu Thr Val Leu
 305 310 315 320
 Arg Glu Ala Leu Arg Ser Ala His Val Glu Ala Gly Ala Val Asp Tyr
 325 330 335
 Val Glu Thr His Gly Thr Gly Thr Ser Leu Gly Asp Pro Ile Glu Val
 340 345 350
 Glu Ala Leu Arg Ala Thr Val Gly Pro Ala Arg Ser Asp Gly Thr Arg
 355 360 365
 Cys Val Leu Gly Ala Val Lys Thr Asn Ile Gly His Leu Glu Ala Ala
 370 375 380
 Ala Gly Val Ala Gly Leu Ile Lys Ala Ala Leu Ser Leu Thr His Glu
 385 390 395 400
 Arg Ile Pro Arg Asn Leu Asn Phe Arg Thr Leu Asn Pro Arg Ile Arg
 405 410 415
 Leu Glu Gly Ser Ala Leu Ala Leu Ala Thr Glu Pro Val Pro Trp Pro
 420 425 430
 Arg Thr Asp Arg Pro Arg Phe Ala Gly Val Ser Ser Phe Gly Met Ser
 435 440 445
 Gly Thr Asn Ala His Val Val Leu Glu Glu Ala Pro Ala Val Glu Leu
 450 455 460
 Trp Pro Ala Ala Pro Glu Arg Ser Ala Glu Leu Leu Val Leu Ser Gly
 465 470 475 480
 Lys Ser Glu Gly Ala Leu Asp Ala Gln Ala Ala Arg Leu Arg Glu His
 485 490 495
 Leu Asp Met His Pro Glu Leu Gly Leu Gly Asp Val Ala Phe Ser Leu
 500 505 510
 Ala Thr Thr Arg Ser Ala Met Ser His Arg Leu Ala Val Ala Val Thr
 515 520 525

Ser Arg Glu Gly Leu Leu Ala Ala Leu Ser Ala Val Ala Gln Gly Gln
 530 535 540
 Thr Pro Ala Gly Ala Ala Arg Cys Ile Ala Ser Ser Ser Arg Gly Lys
 545 550 555 560
 Leu Ala Phe Leu Phe Thr Gly Gln Gly Ala Gln Thr Pro Gly Met Gly
 565 570 575
 Arg Gly Leu Cys Ala Ala Trp Pro Ala Phe Arg Glu Ala Phe Asp Arg
 580 585 590
 Cys Val Ala Leu Phe Asp Arg Glu Leu Asp Arg Pro Leu Arg Glu Val
 595 600 605
 Met Trp Ala Glu Ala Gly Ser Ala Glu Ser Leu Leu Leu Asp Gln Thr
 610 615 620
 Ala Phe Thr Gln Pro Ala Leu Phe Ala Val Glu Tyr Ala Leu Thr Ala
 625 630 635 640
 Leu Trp Arg Ser Trp Gly Val Glu Pro Glu Leu Leu Val Gly His Ser
 645 650 655
 Ile Gly Glu Leu Val Ala Ala Cys Val Ala Gly Val Phe Ser Leu Glu
 660 665 670
 Asp Gly Val Arg Leu Val Ala Ala Arg Gly Arg Leu Met Gln Gly Leu
 675 680 685
 Ser Ala Gly Gly Ala Met Val Ser Leu Gly Ala Pro Glu Ala Glu Val
 690 695 700
 Ala Ala Ala Val Ala Pro His Ala Ala Ser Val Ser Ile Ala Ala Val
 705 710 715 720
 Asn Gly Pro Glu Gln Val Val Ile Ala Gly Val Glu Gln Ala Val Gln
 725 730 735
 Ala Ile Ala Ala Gly Phe Ala Ala Arg Gly Ala Arg Thr Lys Arg Leu
 740 745 750
 His Val Ser His Ala Phe His Ser Pro Leu Met Glu Pro Met Leu Glu
 755 760 765
 Glu Phe Gly Arg Val Ala Ala Ser Val Thr Tyr Arg Arg Pro Ser Val
 770 775 780
 Ser Leu Val Ser Asn Leu Ser Gly Lys Val Val Thr Asp Glu Leu Ser
 785 790 795 800
 Ala Pro Gly Tyr Trp Val Arg His Val Arg Glu Ala Val Arg Phe Ala
 805 810 815
 Asp Gly Val Lys Ala Leu His Glu Ala Gly Ala Gly Thr Phe Val Glu
 820 825 830
 Val Gly Pro Lys Pro Thr Leu Leu Gly Leu Leu Pro Ala Cys Leu Pro
 835 840 845
 Glu Ala Glu Pro Thr Leu Leu Ala Ser Leu Arg Ala Gly Arg Glu Glu
 850 855 860
 Ala Ala Gly Val Leu Glu Ala Leu Gly Arg Leu Trp Ala Ala Gly Gly
 865 870 875 880

Ser Val Ser Trp Pro Gly Val Phe Pro Thr Ala Gly Arg Arg Val Pro
 885 890 895
 Leu Pro Thr Tyr Pro Trp Gln Arg Gln Arg Tyr Trp Ile Glu Ala Pro
 900 905 910
 Ala Glu Gly Leu Gly Ala Thr Ala Ala Asp Ala Leu Ala Gln Trp Phe
 915 920 925
 Tyr Arg Val Asp Trp Pro Glu Met Pro Arg Ser Ser Val Asp Ser Arg
 930 935 940
 Arg Ala Arg Ser Gly Gly Trp Leu Val Leu Ala Asp Arg Gly Gly Val
 945 950 955 960
 Gly Glu Ala Ala Ala Ala Ala Leu Ser Ser Gln Gly Cys Ser Cys Ala
 965 970 975
 Val Leu His Ala Pro Ala Glu Ala Ser Ala Val Ala Glu Gln Val Thr
 980 985 990
 Gln Ala Leu Gly Gly Arg Asn Asp Trp Gln Gly Val Leu Tyr Leu Trp
 995 1000 1005
 Gly Leu Asp Ala Val Val Glu Ala Gly Ala Ser Ala Glu Glu Val Ala
 1010 1015 1020
 Lys Val Thr His Leu Ala Ala Ala Pro Val Leu Ala Leu Ile Gln Ala
 1025 1030 1035 1040
 Leu Gly Thr Gly Pro Arg Ser Pro Arg Leu Trp Ile Val Thr Arg Gly
 1045 1050 1055
 Ala Cys Thr Val Gly Gly Glu Pro Asp Ala Ala Pro Cys Gln Ala Ala
 1060 1065 1070
 Leu Trp Gly Met Gly Arg Val Ala Ala Leu Glu His Pro Gly Ser Trp
 1075 1080 1085
 Gly Gly Leu Val Asp Leu Asp Pro Glu Glu Ser Pro Thr Glu Val Glu
 1090 1095 1100
 Ala Leu Val Ala Glu Leu Leu Ser Pro Asp Ala Glu Asp Gln Leu Ala
 1105 1110 1115 1120
 Phe Arg Gln Gly Arg Arg Arg Ala Ala Arg Leu Val Ala Ala Pro Pro
 1125 1130 1135
 Glu Gly Asn Ala Ala Pro Val Ser Leu Ser Ala Glu Gly Ser Tyr Leu
 1140 1145 1150
 Val Thr Gly Gly Leu Gly Ala Leu Gly Leu Leu Val Ala Arg Trp Leu
 1155 1160 1165
 Val Glu Arg Gly Ala Gly His Leu Val Leu Ile Ser Arg His Gly Leu
 1170 1175 1180
 Pro Asp Arg Glu Glu Trp Gly Arg Asp Gln Pro Pro Glu Val Arg Ala
 1185 1190 1195 1200
 Arg Ile Ala Ala Ile Glu Ala Leu Glu Ala Gln Gly Ala Arg Val Thr
 1205 1210 1215
 Val Ala Ala Val Asp Val Ala Asp Ala Glu Gly Met Ala Ala Leu Leu
 1220 1225 1230

Ala Ala Val Glu Pro Pro Leu Arg Gly Val Val His Ala Ala Gly Leu
 1235 1240 1245
 Leu Asp Asp Gly Leu Leu Ala His Gln Asp Ala Gly Arg Leu Ala Arg
 1250 1255 1260
 Val Leu Arg Pro Lys Val Glu Gly Ala Trp Val Leu His Thr Leu Thr
 1265 1270 1275 1280
 Arg Glu Gln Pro Leu Asp Leu Phe Val Leu Phe Ser Ser Ala Ser Gly
 1285 1290 1295
 Val Phe Gly Ser Ile Gly Gln Gly Ser Tyr Ala Ala Gly Asn Ala Phe
 1300 1305 1310
 Leu Asp Ala Leu Ala Asp Leu Arg Arg Thr Gln Gly Leu Ala Ala Leu
 1315 1320 1325
 Ser Ile Ala Trp Gly Leu Trp Ala Glu Gly Gly Met Gly Ser Gln Ala
 1330 1335 1340
 Gln Arg Arg Glu His Glu Ala Ser Gly Ile Trp Ala Met Pro Thr Ser
 1345 1350 1355 1360
 Arg Ala Leu Ala Ala Met Glu Trp Leu Leu Gly Thr Arg Ala Thr Gln
 1365 1370 1375
 Arg Val Val Ile Gln Met Asp Trp Ala His Ala Gly Ala Ala Pro Arg
 1380 1385 1390
 Asp Ala Ser Arg Gly Arg Phe Trp Asp Arg Leu Val Thr Ala Thr Lys
 1395 1400 1405
 Glu Ala Ser Ser Ser Ala Val Pro Ala Val Glu Arg Trp Arg Asn Ala
 1410 1415 1420
 Ser Val Val Glu Thr Arg Ser Ala Leu Tyr Glu Leu Val Arg Gly Val
 1425 1430 1435 1440
 Val Ala Gly Val Met Gly Phe Thr Asp Gln Gly Thr Leu Asp Val Arg
 1445 1450 1455
 Arg Gly Phe Ala Glu Gln Gly Leu Asp Ser Leu Met Ala Val Glu Ile
 1460 1465 1470
 Arg Lys Arg Leu Gln Gly Glu Leu Gly Met Pro Leu Ser Ala Thr Leu
 1475 1480 1485
 Ala Phe Asp His Pro Thr Val Glu Arg Leu Val Glu Tyr Leu Leu Ser
 1490 1495 1500
 Gln Ala Leu Glu Leu Gln Asp Arg Thr Asp Val Arg Ser Val Arg Leu
 1505 1510 1515 1520
 Pro Ala Thr Glu Asp Pro Ile Ala Ile Val Gly Ala Ala Cys Arg Phe
 1525 1530 1535
 Pro Gly Gly Val Glu Asp Leu Glu Ser Tyr Trp Gln Leu Leu Thr Glu
 1540 1545 1550
 Gly Val Val Val Ser Thr Glu Val Pro Ala Asp Arg Trp Asn Gly Ala
 1555 1560 1565
 Asp Gly Arg Val Pro Gly Ser Gly Glu Ala Gln Arg Gln Thr Tyr Val
 1570 1575 1580

Pro Arg Gly Gly Phe Leu Arg Glu Val Glu Thr Phe Asp Ala Ala Phe
 1585 1590 1595 1600
 Phe His Ile Ser Pro Arg Glu Ala Met Ser Leu Asp Pro Gln Gln Arg
 1605 1610 1615
 Leu Leu Leu Glu Val Ser Trp Glu Ala Ile Glu Arg Ala Gly Gln Asp
 1620 1625 1630
 Pro Ser Ala Leu Arg Glu Ser Pro Thr Gly Val Phe Val Gly Ala Gly
 1635 1640 1645
 Pro Asn Glu Tyr Ala Glu Arg Val Gln Glu Leu Ala Asp Glu Ala Ala
 1650 1655 1660
 Gly Leu Tyr Ser Gly Thr Gly Asn Met Leu Ser Val Ala Ala Gly Arg
 1665 1670 1675 1680
 Leu Ser Phe Phe Leu Gly Leu His Gly Pro Thr Leu Ala Val Asp Thr
 1685 1690 1695
 Ala Cys Ser Ser Ser Leu Val Ala Leu His Leu Gly Cys Gln Ser Leu
 1700 1705 1710
 Arg Arg Gly Glu Cys Asp Gln Ala Leu Val Gly Gly Val Asn Met Leu
 1715 1720 1725
 Leu Ser Pro Lys Thr Phe Ala Leu Leu Ser Arg Met His Ala Leu Ser
 1730 1735 1740
 Pro Gly Gly Arg Cys Lys Thr Phe Ser Ala Asp Ala Asp Gly Tyr Ala
 1745 1750 1755 1760
 Arg Ala Glu Gly Cys Ala Val Val Val Leu Lys Arg Leu Ser Asp Ala
 1765 1770 1775
 Gln Arg Asp Arg Asp Pro Ile Leu Ala Val Ile Arg Gly Thr Ala Ile
 1780 1785 1790
 Asn His Asp Gly Pro Ser Ser Gly Leu Thr Val Pro Ser Gly Pro Ala
 1795 1800 1805
 Gln Glu Ala Leu Leu Arg Gln Ala Leu Ala His Ala Gly Val Val Pro
 1810 1815 1820
 Ala Asp Val Asp Phe Val Glu Cys His Gly Thr Gly Thr Ala Leu Gly
 1825 1830 1835 1840
 Asp Pro Ile Glu Val Arg Ala Leu Ser Asp Val Tyr Gly Gln Ala Arg
 1845 1850 1855
 Pro Ala Asp Arg Pro Leu Ile Leu Gly Ala Ala Lys Ala Asn Leu Gly
 1860 1865 1870
 His Met Glu Pro Ala Ala Gly Leu Ala Gly Leu Leu Lys Ala Val Leu
 1875 1880 1885
 Ala Leu Gly Gln Glu Gln Ile Pro Ala Gln Pro Glu Leu Gly Glu Leu
 1890 1895 1900
 Asn Pro Leu Leu Pro Trp Glu Ala Leu Pro Val Ala Val Ala Arg Ala
 1905 1910 1915 1920
 Ala Val Pro Trp Pro Arg Thr Asp Arg Pro Arg Phe Ala Gly Val Ser
 1925 1930 1935

Ser Phe Gly Met Ser Gly Thr Asn Ala His Val Val Leu Glu Glu Ala
 1940 1945 1950
 Pro Ala Val Glu Leu Trp Pro Ala Ala Pro Glu Arg Ser Ala Glu Leu
 1955 1960 1965
 Leu Val Leu Ser Gly Lys Ser Glu Gly Ala Leu Asp Ala Gln Ala Ala
 1970 1975 1980
 Arg Leu Arg Glu His Leu Asp Met His Pro Glu Leu Gly Leu Gly Asp
 1985 1990 1995 2000
 Val Ala Phe Ser Leu Ala Thr Thr Arg Ser Ala Met Asn His Arg Leu
 2005 2010 2015
 Ala Val Ala Val Thr Ser Arg Glu Gly Leu Leu Ala Ala Leu Ser Ala
 2020 2025 2030
 Val Ala Gln Gly Gln Thr Pro Pro Gly Ala Ala Arg Cys Ile Ala Ser
 2035 2040 2045
 Ser Ser Arg Gly Lys Leu Ala Phe Leu Phe Thr Gly Gln Gly Ala Gln
 2050 2055 2060
 Thr Pro Gly Met Gly Arg Gly Leu Cys Ala Ala Trp Pro Ala Phe Arg
 2065 2070 2075 2080
 Glu Ala Phe Asp Arg Cys Val Ala Leu Phe Asp Arg Glu Leu Asp Arg
 2085 2090 2095
 Pro Leu Arg Glu Val Met Trp Ala Glu Pro Gly Ser Ala Glu Ser Leu
 2100 2105 2110
 Leu Leu Asp Gln Thr Ala Phe Thr Gln Pro Ala Leu Phe Thr Val Glu
 2115 2120 2125
 Tyr Ala Leu Thr Ala Leu Trp Arg Ser Trp Gly Val Glu Pro Glu Leu
 2130 2135 2140
 Val Ala Gly His Ser Ala Gly Glu Leu Val Ala Ala Cys Val Ala Gly
 2145 2150 2155 2160
 Val Phe Ser Leu Glu Asp Gly Val Arg Leu Val Ala Ala Arg Gly Arg
 2165 2170 2175
 Leu Met Gln Gly Leu Ser Ala Gly Gly Ala Met Val Ser Leu Gly Ala
 2180 2185 2190
 Pro Glu Ala Glu Val Ala Ala Ala Val Ala Pro His Ala Ala Ser Val
 2195 2200 2205
 Ser Ile Ala Ala Val Asn Gly Pro Glu Gln Val Val Ile Ala Gly Val
 2210 2215 2220
 Glu Gln Ala Val Gln Ala Ile Ala Ala Gly Phe Ala Ala Arg Gly Ala
 2225 2230 2235 2240
 Arg Thr Lys Arg Leu His Val Ser His Ala Ser His Ser Pro Leu Met
 2245 2250 2255
 Glu Pro Met Leu Glu Glu Phe Gly Arg Val Ala Ala Ser Val Thr Tyr
 2260 2265 2270
 Arg Arg Pro Ser Val Ser Leu Val Ser Asn Leu Ser Gly Lys Val Val
 2275 2280 2285

Ala Asp Glu Leu Ser Ala Pro Gly Tyr Trp Val Arg His Val Arg Glu
 2290 2295 2300
 Ala Val Arg Phe Ala Asp Gly Val Lys Ala Leu His Glu Ala Gly Ala
 2305 2310 2315 2320
 Gly Thr Phe Val Glu Val Gly Pro Lys Pro Thr Leu Leu Gly Leu Leu
 2325 2330 2335
 Pro Ala Cys Leu Pro Glu Ala Glu Pro Thr Leu Leu Ala Ser Leu Arg
 2340 2345 2350
 Ala Gly Arg Glu Glu Ala Ala Gly Val Leu Glu Ala Leu Gly Arg Leu
 2355 2360 2365
 Trp Ala Ala Gly Gly Ser Val Ser Trp Pro Gly Val Phe Pro Thr Ala
 2370 2375 2380
 Gly Arg Arg Val Pro Leu Pro Thr Tyr Pro Trp Gln Arg Gln Arg Tyr
 2385 2390 2395 2400
 Trp Pro Asp Ile Glu Pro Asp Ser Arg Arg His Ala Ala Ala Asp Pro
 2405 2410 2415
 Thr Gln Gly Trp Phe Tyr Arg Val Asp Trp Pro Glu Ile Pro Arg Ser
 2420 2425 2430
 Leu Gln Lys Ser Glu Glu Ala Ser Arg Gly Ser Trp Leu Val Leu Ala
 2435 2440 2445
 Asp Lys Gly Gly Val Gly Glu Ala Val Ala Ala Ala Leu Ser Thr Arg
 2450 2455 2460
 Gly Leu Pro Cys Val Val Leu His Ala Pro Ala Glu Thr Ser Ala Thr
 2465 2470 2475 2480
 Ala Glu Leu Val Thr Glu Ala Ala Gly Gly Arg Ser Asp Trp Gln Val
 2485 2490 2495
 Val Leu Tyr Leu Trp Gly Leu Asp Ala Val Val Gly Ala Glu Ala Ser
 2500 2505 2510
 Ile Asp Glu Ile Gly Asp Ala Thr Arg Arg Ala Thr Ala Pro Val Leu
 2515 2520 2525
 Gly Leu Ala Arg Phe Leu Ser Thr Val Ser Cys Ser Pro Arg Leu Trp
 2530 2535 2540
 Val Val Thr Arg Gly Ala Cys Ile Val Gly Asp Glu Pro Ala Ile Ala
 2545 2550 2555 2560
 Pro Cys Gln Ala Ala Leu Trp Gly Met Gly Arg Val Ala Ala Leu Glu
 2565 2570 2575
 His Pro Gly Ala Trp Gly Gly Leu Val Asp Leu Asp Pro Arg Ala Ser
 2580 2585 2590
 Pro Pro Gln Ala Ser Pro Ile Asp Gly Glu Met Leu Val Thr Glu Leu
 2595 2600 2605
 Leu Ser Gln Glu Thr Glu Asp Gln Leu Ala Phe Arg His Gly Arg Arg
 2610 2615 2620
 His Ala Ala Arg Leu Val Ala Ala Pro Pro Gln Gly Gln Ala Ala Pro
 2625 2630 2635 2640

Val Ser Leu Ser Ala Glu Ala Ser Tyr Leu Val Thr Gly Gly Leu Gly
 2645 2650 2655
 Gly Leu Gly Leu Ile Val Ala Gln Trp Leu Val Glu Leu Gly Ala Arg
 2660 2665 2670
 His Leu Val Leu Thr Ser Arg Arg Gly Leu Pro Asp Arg Gln Ala Trp
 2675 2680 2685
 Cys Glu Gln Gln Pro Pro Glu Ile Arg Ala Arg Ile Ala Ala Val Glu
 2690 2695 2700
 Ala Leu Glu Ala Arg Gly Ala Arg Val Thr Val Ala Ala Val Asp Val
 2705 2710 2715 2720
 Ala Asp Val Glu Pro Met Thr Ala Leu Val Ser Ser Val Glu Pro Pro
 2725 2730 2735
 Leu Arg Gly Val Val His Ala Ala Gly Val Ser Val Met Arg Pro Leu
 2740 2745 2750
 Ala Glu Thr Asp Glu Thr Leu Leu Glu Ser Val Leu Arg Pro Lys Val
 2755 2760 2765
 Ala Gly Ser Trp Leu Leu His Arg Leu Leu His Gly Arg Pro Leu Asp
 2770 2775 2780
 Leu Phe Val Leu Phe Ser Ser Gly Ala Ala Val Trp Gly Ser His Ser
 2785 2790 2795 2800
 Gln Gly Ala Tyr Ala Ala Ala Asn Ala Phe Leu Asp Gly Leu Ala His
 2805 2810 2815
 Leu Arg Arg Ser Gln Ser Leu Pro Ala Leu Ser Val Ala Trp Gly Leu
 2820 2825 2830
 Trp Ala Glu Gly Gly Met Ala Asp Ala Glu Ala His Ala Arg Leu Ser
 2835 2840 2845
 Asp Ile Gly Val Leu Pro Met Ser Thr Ser Ala Ala Leu Ser Ala Leu
 2850 2855 2860
 Gln Arg Leu Val Glu Thr Gly Ala Ala Gln Arg Thr Val Thr Arg Met
 2865 2870 2875 2880
 Asp Trp Ala Arg Phe Ala Pro Val Tyr Thr Ala Arg Gly Arg Arg Asn
 2885 2890 2895
 Leu Leu Ser Ala Leu Val Ala Gly Arg Asp Ile Ile Ala Pro Ser Pro
 2900 2905 2910
 Pro Ala Ala Ala Thr Arg Asn Trp Arg Gly Leu Ser Val Ala Glu Ala
 2915 2920 2925
 Arg Val Ala Leu His Glu Ile Val His Gly Ala Val Ala Arg Val Leu
 2930 2935 2940
 Gly Phe Leu Asp Pro Ser Ala Leu Asp Pro Gly Met Gly Phe Asn Glu
 2945 2950 2955 2960
 Gln Gly Leu Asp Ser Leu Met Ala Val Glu Ile Arg Asn Leu Leu Gln
 2965 2970 2975
 Ala Glu Leu Asp Val Arg Leu Ser Thr Thr Leu Ala Phe Asp His Pro
 2980 2985 2990

Thr Val Gln Arg Leu Val Glu His Leu Leu Val Asp Val Leu Lys Leu
 2995 3000 3005
 Glu Asp Arg Ser Asp Thr Gln His Val Arg Ser Leu Ala Ser Asp Glu
 3010 3015 3020
 Pro Ile Ala Ile Val Gly Ala Ala Cys Arg Phe Pro Gly Gly Val Glu
 3025 3030 3035 3040
 Asp Leu Glu Ser Tyr Trp Gln Leu Leu Ala Glu Gly Val Val Val Ser
 3045 3050 3055
 Ala Glu Val Pro Ala Asp Arg Trp Asp Ala Ala Asp Trp Tyr Asp Pro
 3060 3065 3070
 Asp Pro Glu Ile Pro Gly Arg Thr Tyr Val Thr Lys Gly Ala Phe Leu
 3075 3080 3085
 Arg Asp Leu Gln Arg Leu Asp Ala Thr Phe Phe Arg Ile Ser Pro Arg
 3090 3095 3100
 Glu Ala Met Ser Leu Asp Pro Gln Gln Arg Leu Leu Leu Glu Val Ser
 3105 3110 3115 3120
 Trp Glu Ala Leu Glu Ser Ala Gly Ile Ala Pro Asp Thr Leu Arg Asp
 3125 3130 3135
 Ser Pro Thr Gly Val Phe Val Gly Ala Gly Pro Asn Glu Tyr Tyr Thr
 3140 3145 3150
 Gln Arg Leu Arg Gly Phe Thr Asp Gly Ala Ala Gly Leu Tyr Gly Gly
 3155 3160 3165
 Thr Gly Asn Met Leu Ser Val Thr Ala Gly Arg Leu Ser Phe Phe Leu
 3170 3175 3180
 Gly Leu His Gly Pro Thr Leu Ala Met Asp Thr Ala Cys Ser Ser Ser
 3185 3190 3195 3200
 Leu Val Ala Leu His Leu Ala Cys Gln Ser Leu Arg Leu Gly Glu Cys
 3205 3210 3215
 Asp Gln Ala Leu Val Gly Gly Val Asn Val Leu Leu Ala Pro Glu Thr
 3220 3225 3230
 Phe Val Leu Leu Ser Arg Met Arg Ala Leu Ser Pro Asp Gly Arg Cys
 3235 3240 3245
 Lys Thr Phe Ser Ala Asp Ala Asp Gly Tyr Ala Arg Gly Glu Gly Cys
 3250 3255 3260
 Ala Val Val Val Leu Lys Arg Leu Arg Asp Ala Gln Arg Ala Gly Asp
 3265 3270 3275 3280
 Ser Ile Leu Ala Leu Ile Arg Gly Ser Ala Val Asn His Asp Gly Pro
 3285 3290 3295
 Ser Ser Gly Leu Thr Val Pro Asn Gly Pro Ala Gln Gln Ala Leu Leu
 3300 3305 3310
 Arg Gln Ala Leu Ser Gln Ala Gly Val Ser Pro Val Asp Val Asp Phe
 3315 3320 3325
 Val Glu Cys His Gly Thr Gly Thr Ala Leu Gly Asp Pro Ile Glu Val
 3330 3335 3340

Gln Ala Leu Ser Glu Val Tyr Gly Pro Gly Arg Ser Gly Asp Arg Pro
 3345 3350 3355 3360
 Leu Val Leu Gly Ala Ala Lys Ala Asn Val Ala His Leu Glu Ala Ala
 3365 3370 3375
 Ser Gly Leu Ala Ser Leu Leu Lys Ala Val Leu Ala Leu Arg His Glu
 3380 3385 3390
 Gln Ile Pro Ala Gln Pro Glu Leu Gly Glu Leu Asn Pro His Leu Pro
 3395 3400 3405
 Trp Asn Thr Leu Pro Val Ala Val Pro Arg Lys Ala Val Pro Trp Gly
 3410 3415 3420
 Arg Gly Ala Arg Pro Arg Arg Ala Gly Val Ser Ala Phe Gly Leu Ser
 3425 3430 3435 3440
 Gly Thr Asn Val His Val Val Leu Glu Glu Ala Pro Glu Val Glu Pro
 3445 3450 3455
 Ala Pro Ala Ala Pro Ala Arg Pro Val Glu Leu Val Val Leu Ser Ala
 3460 3465 3470
 Lys Ser Ala Ala Ala Leu Asp Ala Ala Ala Arg Leu Ser Ala His
 3475 3480 3485
 Leu Ser Ala His Pro Glu Leu Ser Leu Gly Asp Val Ala Phe Ser Leu
 3490 3495 3500
 Ala Thr Thr Arg Ser Pro Met Glu His Arg Leu Ala Ile Ala Thr Thr
 3505 3510 3515 3520
 Ser Arg Glu Ala Leu Arg Gly Ala Leu Asp Ala Ala Ala Gln Gln Lys
 3525 3530 3535
 Thr Pro Gln Gly Ala Val Arg Gly Lys Ala Val Ser Ser Arg Gly Lys
 3540 3545 3550
 Leu Ala Phe Leu Phe Thr Gly Gln Gly Ala Gln Met Pro Gly Met Gly
 3555 3560 3565
 Arg Gly Leu Tyr Glu Thr Trp Pro Ala Phe Arg Glu Ala Phe Asp Arg
 3570 3575 3580
 Cys Val Ala Leu Phe Asp Arg Glu Ile Asp Gln Pro Leu Arg Glu Val
 3585 3590 3595 3600
 Met Trp Ala Ala Pro Gly Leu Ala Gln Ala Ala Arg Leu Asp Gln Thr
 3605 3610 3615
 Ala Tyr Ala Gln Pro Ala Leu Phe Ala Leu Glu Tyr Ala Leu Ala Ala
 3620 3625 3630
 Leu Trp Arg Ser Trp Gly Val Glu Pro His Val Leu Leu Gly His Ser
 3635 3640 3645
 Ile Gly Glu Leu Val Ala Ala Cys Val Ala Gly Val Phe Ser Leu Glu
 3650 3655 3660
 Asp Ala Val Arg Leu Val Ala Ala Arg Gly Arg Leu Met Gln Ala Leu
 3665 3670 3675 3680
 Pro Ala Gly Gly Ala Met Val Ala Ile Ala Ala Ser Glu Ala Glu Val
 3685 3690 3695

Ala Ala Ser Val Ala Pro His Ala Ala Thr Val Ser Ile Ala Ala Val
 3700 3705 3710
 Asn Gly Pro Asp Ala Val Val Ile Ala Gly Ala Glu Val Gln Val Leu
 3715 3720 3725
 Ala Leu Gly Ala Thr Phe Ala Ala Arg Gly Ile Arg Thr Lys Arg Leu
 3730 3735 3740
 Ala Val Ser His Ala Phe His Ser Pro Leu Met Asp Pro Met Leu Glu
 3745 3750 3755 3760
 Asp Phe Gln Arg Val Ala Ala Thr Ile Ala Tyr Arg Ala Pro Asp Arg
 3765 3770 3775
 Pro Val Val Ser Asn Val Thr Gly His Val Ala Gly Pro Glu Ile Ala
 3780 3785 3790
 Thr Pro Glu Tyr Trp Val Arg His Val Arg Ser Ala Val Arg Phe Gly
 3795 3800 3805
 Asp Gly Ala Lys Ala Leu His Ala Ala Gly Ala Ala Thr Phe Val Glu
 3810 3815 3820
 Val Gly Pro Lys Pro Val Leu Leu Gly Leu Leu Pro Ala Cys Leu Gly
 3825 3830 3835 3840
 Glu Ala Asp Ala Val Leu Val Pro Ser Leu Arg Ala Asp Arg Ser Glu
 3845 3850 3855
 Cys Glu Val Val Leu Ala Ala Leu Gly Ala Trp Tyr Ala Trp Gly Gly
 3860 3865 3870
 Ala Leu Asp Trp Lys Gly Val Phe Pro Asp Gly Ala Arg Arg Val Ala
 3875 3880 3885
 Leu Pro Met Tyr Pro Trp Gln Arg Glu Arg His Trp Met Asp Leu Thr
 3890 3895 3900
 Pro Arg Ser Ala Ala Pro Ala Gly Ile Ala Gly Arg Trp Pro Leu Ala
 3905 3910 3915 3920
 Gly Val Gly Leu Cys Met Pro Gly Ala Val Leu His His Val Leu Ser
 3925 3930 3935
 Ile Gly Pro Arg His Gln Pro Phe Leu Gly Asp His Leu Val Phe Gly
 3940 3945 3950
 Lys Val Val Val Pro Gly Ala Phe His Val Ala Val Ile Leu Ser Ile
 3955 3960 3965
 Ala Ala Glu Arg Trp Pro Glu Arg Ala Ile Glu Leu Thr Gly Val Glu
 3970 3975 3980
 Phe Leu Lys Ala Ile Ala Met Glu Pro Asp Gln Glu Val Glu Leu His
 3985 3990 3995 4000
 Ala Val Leu Thr Pro Glu Ala Ala Gly Asp Gly Tyr Leu Phe Glu Leu
 4005 4010 4015
 Ala Thr Leu Ala Ala Pro Glu Thr Glu Arg Arg Trp Thr Thr His Ala
 4020 4025 4030
 Arg Gly Arg Val Gln Pro Thr Asp Gly Ala Pro Gly Ala Leu Pro Arg
 4035 4040 4045

Leu Glu Val Leu Glu Asp Arg Ala Ile Gln Pro Leu Asp Phe Ala Gly
 4050 4055 4060
 Phe Leu Asp Arg Leu Ser Ala Val Arg Ile Gly Trp Gly Pro Leu Trp
 4065 4070 4075 4080
 Arg Trp Leu Gln Asp Gly Arg Val Gly Asp Glu Ala Ser Leu Ala Thr
 4085 4090 4095
 Leu Val Pro Thr Tyr Pro Asn Ala His Asp Val Ala Pro Leu His Pro
 4100 4105 4110
 Ile Leu Leu Asp Asn Gly Phe Ala Val Ser Leu Leu Ser Thr Arg Ser
 4115 4120 4125
 Glu Pro Glu Asp Asp Gly Thr Pro Pro Leu Pro Phe Ala Val Glu Arg
 4130 4135 4140
 Val Arg Trp Trp Arg Ala Pro Val Gly Arg Val Arg Cys Gly Gly Val
 4145 4150 4155 4160
 Pro Arg Ser Gln Ala Phe Gly Val Ser Ser Phe Val Leu Val Asp Glu
 4165 4170 4175
 Thr Gly Glu Val Val Ala Glu Val Glu Gly Phe Val Cys Arg Arg Ala
 4180 4185 4190
 Pro Arg Glu Val Phe Leu Arg Gln Glu Ser Gly Ala Ser Thr Ala Ala
 4195 4200 4205
 Leu Tyr Arg Leu Asp Trp Pro Glu Ala Pro Leu Pro Asp Ala Pro Ala
 4210 4215 4220
 Glu Arg Ile Glu Glu Ser Trp Val Val Val Ala Ala Pro Gly Ser Glu
 4225 4230 4235 4240
 Met Ala Ala Ala Leu Ala Thr Arg Leu Asn Arg Cys Val Leu Ala Glu
 4245 4250 4255
 Pro Lys Gly Leu Glu Ala Ala Leu Ala Gly Val Ser Pro Ala Gly Val
 4260 4265 4270
 Ile Cys Leu Trp Glu Ala Gly Ala His Glu Glu Ala Pro Ala Ala Ala
 4275 4280 4285
 Gln Arg Val Ala Thr Glu Gly Leu Ser Val Val Gln Ala Leu Arg Asp
 4290 4295 4300
 Arg Ala Val Arg Leu Trp Trp Val Thr Met Gly Ala Val Ala Val Glu
 4305 4310 4315 4320
 Ala Gly Glu Arg Val Gln Val Ala Thr Ala Pro Val Trp Gly Leu Gly
 4325 4330 4335
 Arg Thr Val Met Gln Glu Arg Pro Glu Leu Ser Cys Thr Leu Val Asp
 4340 4345 4350
 Leu Glu Pro Glu Ala Asp Ala Ala Arg Ser Ala Asp Val Leu Leu Arg
 4355 4360 4365
 Glu Leu Gly Arg Ala Asp Asp Glu Thr Gln Val Ala Phe Arg Ser Gly
 4370 4375 4380
 Lys Arg Arg Val Ala Arg Leu Val Lys Ala Thr Thr Pro Glu Gly Leu
 4385 4390 4395 4400

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Leu Val Pro Asp Ala Glu Ser Tyr Arg Leu Glu Ala Gly Gln Lys Gly
 4405 4410 4415
 Thr Leu Asp Gln Leu Arg Leu Ala Pro Ala Gln Arg Arg Ala Pro Gly
 4420 4425 4430
 Pro Gly Glu Val Glu Ile Lys Val Thr Ala Ser Gly Leu Asn Phe Arg
 4435 4440 4445
 Thr Val Leu Ala Val Leu Gly Met Tyr Pro Gly Asp Ala Gly Pro Met
 4450 4455 4460
 Gly Gly Asp Cys Ala Gly Val Ala Thr Ala Val Gly Gln Gly Val Arg
 4465 4470 4475 4480
 His Val Ala Val Gly Asp Ala Val Met Thr Leu Gly Thr Leu His Arg
 4485 4490 4495
 Phe Val Thr Val Asp Ala Arg Leu Val Val Arg Gln Pro Ala Gly Leu
 4500 4505 4510
 Thr Pro Ala Gln Ala Ala Thr Val Pro Val Ala Phe Leu Thr Ala Trp
 4515 4520 4525
 Leu Ala Leu His Asp Leu Gly Asn Leu Arg Arg Gly Glu Arg Val Leu
 4530 4535 4540
 Ile His Ala Ala Ala Gly Gly Val Gly Met Ala Ala Val Gln Ile Ala
 4545 4550 4555 4560
 Arg Trp Ile Gly Ala Glu Val Phe Ala Thr Ala Ser Pro Ser Lys Trp
 4565 4570 4575
 Ala Ala Val Gln Ala Met Gly Val Pro Arg Thr His Ile Ala Ser Ser
 4580 4585 4590
 Arg Thr Leu Glu Phe Ala Glu Thr Phe Arg Gln Val Thr Gly Gly Arg
 4595 4600 4605
 Gly Val Asp Val Val Leu Asn Ala Leu Ala Gly Glu Phe Val Asp Ala
 4610 4615 4620
 Ser Leu Ser Leu Leu Ser Thr Gly Gly Arg Phe Leu Glu Met Gly Lys
 4625 4630 4635 4640
 Thr Asp Ile Arg Asp Arg Ala Ala Val Ala Ala Ala His Pro Gly Val
 4645 4650 4655
 Arg Tyr Arg Val Phe Asp Ile Leu Glu Leu Ala Pro Asp Arg Thr Arg
 4660 4665 4670
 Glu Ile Leu Glu Arg Val Val Glu Gly Phe Ala Ala Gly His Leu Arg
 4675 4680 4685
 Ala Leu Pro Val His Ala Phe Ala Ile Thr Lys Ala Glu Ala Ala Phe
 4690 4695 4700
 Arg Phe Met Ala Gln Ala Arg His Gln Gly Lys Val Val Leu Leu Pro
 4705 4710 4715 4720
 Ala Pro Ser Ala Ala Pro Leu Ala Pro Thr Gly Thr Val Leu Leu Thr
 4725 4730 4735
 Gly Gly Leu Gly Ala Leu Gly Leu His Val Ala Arg Trp Leu Ala Gln
 4740 4745 4750

Gln Gly Val Pro His Met Val Leu Thr Gly Arg Arg Gly Leu Asp Thr
 4755 4760 4765
 Pro Gly Ala Ala Lys Ala Val Ala Glu Ile Glu Ala Leu Gly Ala Arg
 4770 4775 4780
 Val Thr Ile Ala Ala Ser Asp Val Ala Asp Arg Asn Ala Leu Glu Ala
 4785 4790 4795 4800
 Val Leu Gln Ala Ile Pro Ala Glu Trp Pro Leu Gln Gly Val Ile His
 4805 4810 4815
 Ala Ala Gly Ala Leu Asp Asp Gly Val Leu Asp Glu Gln Thr Thr Asp
 4820 4825 4830
 Arg Phe Ser Arg Val Leu Ala Pro Lys Val Thr Gly Ala Trp Asn Leu
 4835 4840 4845
 His Glu Leu Thr Ala Gly Asn Asp Leu Ala Phe Phe Val Leu Phe Ser
 4850 4855 4860
 Ser Met Ser Gly Leu Leu Gly Ser Ala Gly Gln Ser Asn Tyr Ala Ala
 4865 4870 4875 4880
 Ala Asn Thr Phe Leu Asp Ala Leu Ala Ala His Arg Arg Ala Glu Gly
 4885 4890 4895
 Leu Ala Ala Gln Ser Leu Ala Trp Gly Pro Trp Ser Asp Gly Gly Met
 4900 4905 4910
 Ala Ala Gly Leu Ser Ala Ala Leu Gln Ala Arg Leu Ala Arg His Gly
 4915 4920 4925
 Met Gly Ala Leu Ser Pro Ala Gln Gly Thr Ala Leu Leu Gly Gln Ala
 4930 4935 4940
 Leu Ala Arg Pro Glu Thr Gln Leu Gly Ala Met Ser Leu Asp Val Arg
 4945 4950 4955 4960
 Ala Ala Ser Gln Ala Ser Gly Ala Ala Val Pro Pro Val Trp Arg Ala
 4965 4970 4975
 Leu Val Arg Ala Glu Ala Arg His Thr Ala Ala Gly Ala Gln Gly Ala
 4980 4985 4990
 Leu Ala Ala Arg Leu Gly Ala Leu Pro Glu Ala Arg Arg Ala Asp Glu
 4995 5000 5005
 Val Arg Lys Val Val Gln Ala Glu Ile Ala Arg Val Leu Ser Trp Ser
 5010 5015 5020
 Ala Ala Ser Ala Val Pro Val Asp Arg Pro Leu Ser Asp Leu Gly Leu
 5025 5030 5035 5040
 Asp Ser Leu Thr Ala Val Glu Leu Arg Asn Val Leu Gly Gln Arg Val
 5045 5050 5055
 Gly Ala Thr Leu Pro Ala Thr Leu Ala Phe Asp His Pro Thr Val Asp
 5060 5065 5070
 Ala Leu Thr Arg Trp Leu Leu Asp Lys Val Leu Ala Val Ala Glu Pro
 5075 5080 5085
 Ser Val Ser Ser Ala Lys Ser Ser Pro Gln Val Ala Leu Asp Glu Pro
 5090 5095 5100

Ile Ala Ile Ile Gly Ile Gly Cys Arg Phe Pro Gly Gly Val Ala Asp
 5105 5110 5115 5120
 Pro Glu Ser Phe Trp Arg Leu Leu Glu Glu Gly Ser Asp Ala Val Val
 5125 5130 5135
 Glu Val Pro His Glu Arg Trp Asp Ile Asp Ala Phe Tyr Asp Pro Asp
 5140 5145 5150
 Pro Asp Val Arg Gly Lys Met Thr Thr Arg Phe Gly Gly Phe Leu Ser
 5155 5160 5165
 Asp Ile Asp Arg Phe Asp Pro Ala Phe Phe Gly Ile Ser Pro Arg Glu
 5170 5175 5180
 Ala Thr Thr Met Asp Pro Gln Gln Arg Leu Leu Leu Glu Thr Ser Trp
 5185 5190 5195 5200
 Glu Ala Phe Glu Arg Ala Gly Ile Leu Pro Glu Arg Leu Met Gly Ser
 5205 5210 5215
 Asp Thr Gly Val Phe Val Gly Leu Phe Tyr Gln Glu Tyr Ala Ala Leu
 5220 5225 5230
 Ala Gly Gly Ile Glu Ala Phe Asp Gly Tyr Leu Gly Thr Gly Thr Thr
 5235 5240 5245
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 5250 5255 5260
 Pro Ser Leu Thr Val Asp Thr Ala Cys Ser Ser Ser Leu Val Ala Val
 5265 5270 5275 5280
 His Leu Ala Cys Gln Ala Leu Arg Arg Gly Glu Cys Ser Val Ala Leu
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 Ala Gly Gly Val Ala Leu Met Leu Thr Pro Ala Thr Phe Val Glu Phe
 5300 5305 5310
 Ser Arg Leu Arg Gly Leu Ala Pro Asp Gly Arg Cys Lys Ser Phe Ser
 5315 5320 5325
 Ala Ala Ala Asp Gly Val Gly Trp Ser Glu Gly Cys Ala Met Leu Leu
 5330 5335 5340
 Leu Lys Pro Leu Arg Asp Ala Gln Arg Asp Gly Asp Pro Ile Leu Ala
 5345 5350 5355 5360
 Val Ile Arg Gly Thr Ala Val Asn Gln Asp Gly Arg Ser Asn Gly Leu
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 Thr Ala Pro Asn Gly Ser Ser Gln Gln Glu Val Ile Arg Arg Ala Leu
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 Glu Gln Ala Gly Leu Ala Pro Ala Asp Val Ser Tyr Val Glu Cys His
 5395 5400 5405
 Gly Thr Gly Thr Thr Leu Gly Asp Pro Ile Glu Val Gln Ala Leu Gly
 5410 5415 5420
 Ala Val Leu Ala Gln Gly Arg Pro Ser Asp Arg Pro Leu Val Ile Gly
 5425 5430 5435 5440
 Ser Val Lys Ser Asn Ile Gly His Thr Gln Ala Ala Ala Gly Val Ala
 5445 5450 5455

Gly Val Ile Lys Val Ala Leu Ala Leu Glu Arg Gly Leu Ile Pro Arg
 5460 5465 5470
 Ser Leu His Phe Asp Ala Pro Asn Pro His Ile Pro Trp Ser Glu Leu
 5475 5480 5485
 Ala Val Gln Val Ala Ala Lys Pro Val Glu Trp Thr Arg Asn Gly Val
 5490 5495 5500
 Pro Arg Arg Ala Gly Val Ser Ser Phe Gly Val Ser Gly Thr Asn Ala
 5505 5510 5515 5520
 His Val Val Leu Glu Glu Ala Pro Ala Ala Ala Phe Ala Pro Ala Ala
 5525 5530 5535
 Ala Arg Ser Ala Glu Leu Phe Val Leu Ser Ala Lys Ser Ala Ala Ala
 5540 5545 5550
 Leu Asp Ala Gln Ala Ala Arg Leu Ser Ala His Val Val Ala His Pro
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 Glu Leu Gly Leu Gly Asp Leu Ala Phe Ser Leu Ala Thr Thr Arg Ser
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 Pro Met Thr Tyr Arg Leu Ala Val Ala Ala Thr Ser Arg Glu Ala Leu
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 Ser Ala Ala Leu Asp Thr Ala Ala Gln Gly Gln Ala Pro Pro Ala Ala
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 Ala Arg Gly His Ala Ser Thr Gly Ser Ala Pro Lys Val Val Phe Val
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 Phe Pro Gly Gln Gly Ser Gln Trp Leu Gly Met Gly Gln Lys Leu Leu
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 Ser Glu Glu Pro Val Phe Arg Asp Ala Leu Ser Ala Cys Asp Arg Ala
 5650 5655 5660
 Ile Gln Ala Glu Ala Gly Trp Ser Leu Leu Ala Glu Leu Ala Ala Asp
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 Glu Thr Thr Ser Gln Leu Gly Arg Ile Asp Val Val Gln Pro Ala Leu
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 Phe Ala Ile Glu Val Ala Leu Ser Ala Leu Trp Arg Ser Trp Gly Val
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 Glu Pro Asp Ala Val Val Gly His Ser Met Gly Glu Val Ala Ala Ala
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 His Val Ala Gly Ala Leu Ser Leu Glu Asp Ala Val Ala Ile Ile Cys
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 Arg Arg Ser Leu Leu Leu Arg Arg Ile Ser Gly Gln Gly Glu Met Ala
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 Val Val Glu Leu Ser Leu Ala Glu Ala Glu Ala Ala Leu Leu Gly Tyr
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 Glu Asp Arg Leu Ser Val Ala Val Ser Asn Ser Pro Arg Ser Thr Val
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 Leu Ala Gly Glu Pro Ala Ala Leu Ala Glu Val Leu Ala Ile Leu Ala
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Ala Lys Gly Val Phe Cys Arg Arg Val Lys Val Asp Val Ala Ser His
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 Ser Pro Gln Ile Asp Pro Leu Arg Asp Glu Leu Leu Ala Ala Leu Gly
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 Glu Leu Glu Pro Arg Gln Ala Thr Val Ser Met Arg Ser Thr Val Thr
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 Ser Thr Ile Met Ala Gly Pro Glu Leu Val Ala Ser Tyr Trp Ala Asp
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 Asn Val Arg Gln Pro Val Arg Phe Ala Glu Ala Val Gln Ser Leu Met
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 Glu Asp Gly His Gly Leu Phe Val Glu Met Ser Pro His Pro Ile Leu
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 Thr Thr Ser Val Glu Glu Ile Arg Arg Ala Thr Lys Arg Glu Gly Val
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 Ala Val Gly Ser Leu Arg Arg Gly Gln Asp Glu Arg Leu Ser Met Leu
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 Glu Ala Leu Gly Ala Leu Trp Val His Gly Gln Ala Val Gly Trp Glu
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 Arg Leu Phe Ser Ala Gly Gly Ala Gly Leu Arg Arg Val Pro Leu Pro
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 Thr Tyr Pro Trp Gln Arg Glu Arg Tyr Trp Val Asp Ala Pro Thr Gly
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 Gly Ala Ala Gly Gly Ser Arg Phe Ala His Ala Gly Ser His Pro Leu
 5985 5990 5995 6000
 Leu Gly Glu Met Gln Thr Leu Ser Thr Gln Arg Ser Thr Arg Val Trp
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 Glu Thr Thr Leu Asp Leu Lys Arg Leu Pro Trp Leu Gly Asp His Arg
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 His Val Ala Ser Arg Val Pro Gly His Gly Gly Ala Ala Phe Arg Ser
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 His Ala Arg Gly Val Leu Arg Gln Ile Glu Arg Ala Glu Val Pro Ala
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 Arg Leu Asp Leu Ala Ala Leu Arg Ala Arg Leu Gln Ala Ser Ala Pro
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 Ala Ala Ala Thr Tyr Ala Ala Leu Ala Glu Met Gly Leu Glu Tyr Gly
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Pro Ala Phe Gln Gly Leu Val Glu Leu Trp Arg Gly Glu Gly Glu Ala
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 Leu Gly Arg Val Arg Leu Pro Glu Ala Ala Gly Ser Pro Ala Ala Cys
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 Phe Ala Asp Arg Gly Glu Ala Thr Pro Trp Val Pro Val Glu Ile Gly
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 Ser Leu Arg Trp Phe Gln Arg Pro Ser Gly Glu Leu Trp Cys His Ala
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 Arg Ser Val Ser His Gly Lys Pro Thr Pro Asp Arg Arg Ser Thr Asp
 6245 6250 6255
 Phe Trp Val Val Asp Ser Thr Gly Ala Ile Val Ala Glu Ile Ser Gly
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 Leu Val Ala Gln Arg Leu Ala Gly Gly Val Arg Arg Arg Glu Glu Asp
 6275 6280 6285
 Asp Trp Phe Met Glu Pro Ala Trp Glu Pro Thr Ala Val Pro Gly Ser
 6290 6295 6300
 Glu Val Met Ala Gly Arg Trp Leu Leu Ile Gly Ser Gly Gly Gly Leu
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 Gly Ala Ala Leu His Ser Ala Leu Thr Glu Ala Gly His Ser Val Val
 6325 6330 6335
 His Ala Thr Gly Arg Gly Thr Ser Ala Ala Gly Leu Gln Ala Leu Leu
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 Asp Ala Leu Glu Glu Ser Leu Val Arg Gly Cys Asp Ser Val Leu Trp
 6385 6390 6395 6400
 Thr Val Gln Ala Val Ala Gly Ala Gly Phe Arg Asp Pro Pro Arg Leu
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 Trp Leu Val Thr Arg Gly Ala Gln Ala Ile Gly Ala Gly Asp Val Ser
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 Val Ala Gln Ala Pro Leu Leu Gly Leu Gly Arg Val Ile Ala Leu Glu
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 His Ala Glu Leu Arg Cys Ala Arg Ile Asp Leu Asp Pro Ala Arg Arg
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 Asp Gly Glu Val Asp Glu Leu Leu Ala Glu Leu Leu Ala Asp Asp Ala
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 Glu Glu Glu Val Ala Phe Arg Gly Gly Glu Arg Arg Val Ala Arg Leu
 6485 6490 6495
 Val Arg Arg Leu Pro Glu Thr Asp Cys Arg Glu Lys Ile Glu Pro Ala
 6500 6505 6510

Glu Gly Arg Pro Phe Arg Leu Glu Ile Asp Gly Ser Gly Val Leu Asp
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 Asp Leu Val Leu Arg Ala Thr Glu Arg Arg Pro Pro Gly Pro Gly Glu
 6530 6535 6540
 Val Glu Ile Ala Val Glu Ala Ala Gly Leu Asn Phe Leu Asp Val Met
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 Arg Ala Met Gly Ile Tyr Pro Gly Pro Gly Asp Gly Pro Val Ala Leu
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 Ser Leu Arg Ile Gly Gln Asp Val Val Ala Val Ala Pro Phe Ser Phe
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 Gly Thr His Val Thr Ile Asp Ala Arg Met Leu Ala Pro Arg Pro Ala
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 Ala Trp Tyr Gly Leu Val His Leu Gly Arg Leu Arg Ala Gly Glu Arg
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 Val Leu Ile His Ser Ala Thr Gly Gly Thr Gly Leu Ala Ala Val Gln
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 Ile Ala Arg His Leu Gly Ala Glu Ile Phe Ala Thr Ala Gly Thr Pro
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 Glu Lys Arg Ala Trp Leu Arg Glu Gln Gly Ile Ala His Val Met Asp
 6690 6695 6700
 Ser Arg Ser Leu Asp Phe Ala Glu Gln Val Leu Ala Ala Thr Lys Gly
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 Glu Gly Val Asp Val Val Leu Asn Ser Leu Ser Gly Ala Ala Ile Asp
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 Ala Ser Leu Ser Thr Leu Val Pro Asp Gly Arg Phe Ile Glu Leu Gly
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 Lys Thr Asp Ile Tyr Ala Asp Arg Ser Leu Gly Leu Ala His Phe Arg
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 Lys Ser Leu Ser Tyr Ser Ala Val Asp Leu Ala Gly Leu Ala Val Arg
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 Arg Pro Glu Arg Val Ala Ala Leu Leu Ala Glu Val Val Asp Leu Leu
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 Ala Arg Gly Ala Leu Gln Pro Leu Pro Val Glu Ile Phe Pro Leu Ser
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 Arg Ala Ala Asp Ala Phe Arg Lys Met Ala Gln Ala Gln His Leu Gly
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 Lys Leu Val Leu Ala Leu Glu Asp Pro Asp Val Arg Ile Arg Val Pro
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 Gly Glu Ser Gly Val Ala Ile Arg Ala Asp Gly Ala Tyr Leu Val Thr
 6850 6855 6860

Gly Gly Leu Gly Gly Leu Gly Leu Ser Val Ala Gly Trp Leu Ala Glu
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 Gln Gly Ala Gly His Leu Val Leu Val Gly Arg Ser Gly Ala Val Ser
 6885 6890 6895
 Ala Glu Gln Gln Thr Ala Val Ala Ala Leu Glu Ala His Gly Ala Arg
 6900 6905 6910
 Val Thr Val Ala Arg Ala Asp Val Ala Asp Arg Ala Gln Met Glu Arg
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 Ile Leu Arg Glu Val Thr Ala Ser Gly Met Pro Leu Arg Gly Val Val
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 His Ala Ala Gly Ile Leu Asp Asp Gly Leu Leu Met Gln Gln Thr Pro
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 Ala Arg Phe Arg Ala Val Met Ala Pro Lys Val Arg Gly Ala Leu His
 6965 6970 6975
 Leu His Ala Leu Thr Arg Glu Ala Pro Leu Ser Phe Phe Val Leu Tyr
 6980 6985 6990
 Ala Ser Gly Ala Gly Leu Leu Gly Ser Pro Gly Gln Gly Asn Tyr Ala
 6995 7000 7005
 Ala Ala Asn Thr Phe Leu Asp Ala Leu Ala His His Arg Arg Ala Gln
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 Gly Leu Pro Ala Leu Ser Ile Asp Trp Gly Leu Phe Ala Asp Val Gly
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 Thr Arg Ser Leu Thr Pro Asp Glu Gly Leu Trp Ala Leu Glu Arg Leu
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 Leu Asp Gly Asp Arg Thr Gln Ala Gly Val Met Pro Phe Asp Val Arg
 7075 7080 7085
 Gln Trp Val Glu Phe Tyr Pro Ala Ala Ala Ser Ser Arg Arg Leu Ser
 7090 7095 7100
 Arg Leu Met Thr Ala Arg Arg Val Ala Ser Gly Arg Leu Ala Gly Asp
 7105 7110 7115 7120
 Arg Asp Leu Leu Glu Arg Leu Ala Thr Ala Glu Ala Gly Ala Arg Ala
 7125 7130 7135
 Gly Met Leu Gln Glu Val Val Arg Ala Gln Val Ser Gln Val Leu Arg
 7140 7145 7150
 Leu Ser Glu Gly Lys Leu Asp Val Asp Ala Pro Leu Thr Ser Leu Gly
 7155 7160 7165
 Met Asp Ser Leu Met Gly Leu Glu Leu Arg Asn Arg Ile Glu Ala Val
 7170 7175 7180
 Leu Gly Ile Thr Met Pro Ala Thr Leu Leu Trp Thr Tyr Pro Thr Val
 7185 7190 7195 7200
 Ala Ala Leu Ser Ala His Leu Ala Ser His Val Val Ser Thr Gly Asp
 7205 7210 7215

Gly Glu Ser Ala Arg Pro Pro Asp Thr Gly Ser Val Ala Pro Thr Thr
7220 7225 7230

His Glu Val Ala Ser Leu Asp Glu Asp Gly Leu Phe Ala Leu Ile Asp
7235 7240 7245

Glu Ser Leu Ala Arg Ala Gly Lys Arg
7250 7255

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<211> 3798

<212> PRT

<213> Sorangium cellulosum

<400> 6

Val Thr Asp Arg Glu Gly Gln Leu Leu Glu Arg Leu Arg Glu Val Thr
1 5 10 15

Leu Ala Leu Arg Lys Thr Leu Asn Glu Arg Asp Thr Leu Glu Leu Glu
20 25 30

Lys Thr Glu Pro Ile Ala Ile Val Gly Ile Gly Cys Arg Phe Pro Gly
35 40 45

Gly Ala Gly Thr Pro Glu Ala Phe Trp Glu Leu Leu Asp Asp Gly Arg
50 55 60

Asp Ala Ile Arg Pro Leu Glu Glu Arg Trp Ala Leu Val Gly Val Asp
65 70 75 80

Pro Gly Asp Asp Val Pro Arg Trp Ala Gly Leu Leu Thr Glu Ala Ile
85 90 95

Asp Gly Phe Asp Ala Ala Phe Phe Gly Ile Ala Pro Arg Glu Ala Arg
100 105 110

Ser Leu Asp Pro Gln His Arg Leu Leu Leu Glu Val Ala Trp Glu Gly
115 120 125

Phe Glu Asp Ala Gly Ile Pro Pro Arg Ser Leu Val Gly Ser Arg Thr
130 135 140

Gly Val Phe Val Gly Val Cys Ala Thr Glu Tyr Leu His Ala Ala Val
145 150 155 160

Ala His Gln Pro Arg Glu Glu Arg Asp Ala Tyr Ser Thr Thr Gly Asn
165 170 175

Met Leu Ser Ile Ala Ala Gly Arg Leu Ser Tyr Thr Leu Gly Leu Gln
180 185 190

Gly Pro Cys Leu Thr Val Asp Thr Ala Cys Ser Ser Ser Leu Val Ala
195 200 205

Ile His Leu Ala Cys Arg Ser Leu Arg Ala Arg Glu Ser Asp Leu Ala
210 215 220

Leu Ala Gly Gly Val Asn Met Leu Leu Ser Pro Asp Thr Met Arg Ala
225 230 235 240

Leu Ala Arg Thr Gln Ala Leu Ser Pro Asn Gly Arg Cys Gln Thr Phe
245 250 255

Asp Ala Ser Ala Asn Gly Phe Val Arg Gly Glu Gly Cys Gly Leu Ile

[illegible]

610					615					620					
Val 625	Leu	Phe	Ala	Met	Glu 630	Val	Ala	Leu	Ser	Ala 635	Leu	Trp	Arg	Ser	Trp 640
Gly	Val	Glu	Pro	Glu 645	Ala	Val	Val	Gly	His 650	Ser	Met	Gly	Glu	Val	Ala 655
Ala	Ala	His	Val 660	Ala	Gly	Ala	Leu	Ser 665	Leu	Glu	Asp	Ala	Val 670	Ala	Ile
Ile	Cys	Arg 675	Arg	Ser	Arg	Leu	Leu 680	Arg	Arg	Ile	Ser	Gly 685	Gln	Gly	Glu
Met	Ala 690	Leu	Val	Glu	Leu	Ser 695	Leu	Glu	Glu	Ala	Glu 700	Ala	Ala	Leu	Arg
Gly 705	His	Glu	Gly	Arg	Leu 710	Ser	Val	Ala	Val	Ser 715	Asn	Ser	Pro	Arg	Ser 720
Thr	Val	Leu	Ala	Gly 725	Glu	Pro	Ala	Ala	Leu 730	Ser	Glu	Val	Leu	Ala 735	Ala
Leu	Thr	Ala	Lys 740	Gly	Val	Phe	Trp	Arg 745	Gln	Val	Lys	Val	Asp 750	Val	Ala
Ser	His	Ser 755	Pro	Gln	Val	Asp	Pro 760	Leu	Arg	Glu	Glu	Leu 765	Ile	Ala	Ala
Leu	Gly 770	Ala	Ile	Arg	Pro	Arg 775	Ala	Ala	Ala	Val	Pro 780	Met	Arg	Ser	Thr
Val 785	Thr	Gly	Gly	Val	Ile 790	Ala	Gly	Pro	Glu	Leu 795	Gly	Ala	Ser	Tyr	Trp 800
Ala	Asp	Asn	Leu	Arg 805	Gln	Pro	Val	Arg	Phe 810	Ala	Ala	Ala	Ala	Gln 815	Ala
Leu	Leu	Glu	Gly 820	Gly	Pro	Ala	Leu	Phe 825	Ile	Glu	Met	Ser	Pro 830	His	Pro
Ile	Leu	Val 835	Pro	Pro	Leu	Asp	Glu 840	Ile	Gln	Thr	Ala	Ala 845	Glu	Gln	Gly
Gly	Ala 850	Ala	Val	Gly	Ser	Leu 855	Arg	Arg	Gly	Gln	Asp 860	Glu	Arg	Ala	Thr
Leu 865	Leu	Glu	Ala	Leu	Gly 870	Thr	Leu	Trp	Ala	Ser 875	Gly	Tyr	Pro	Val	Ser 880
Trp	Ala	Arg	Leu	Phe 885	Pro	Ala	Gly	Gly	Arg 890	Arg	Val	Pro	Leu	Pro 895	Thr
Tyr	Pro	Trp	Gln 900	His	Glu	Arg	Cys	Trp 905	Ile	Glu	Val	Glu	Pro 910	Asp	Ala
Arg	Arg	Leu 915	Ala	Ala	Ala	Asp	Pro 920	Thr	Lys	Asp	Trp	Phe 925	Tyr	Arg	Thr
Asp	Trp 930	Pro	Glu	Val	Pro	Arg 935	Ala	Ala	Pro	Lys	Ser 940	Glu	Thr	Ala	His
Gly 945	Ser	Trp	Leu	Leu	Leu 950	Ala	Asp	Arg	Gly	Gly 955	Val	Gly	Glu	Ala	Val 960
Ala	Ala	Ala	Leu	Ser	Thr	Arg	Gly	Leu	Ser	Cys	Thr	Val	Leu	His	Ala

9

1315	1320	1325
Gly Leu Trp Ala Glu Gly Gly Met Val Asp Ala Lys Ala His Ala Arg 1330	1335	1340
Leu Ser Asp Ile Gly Val Leu Pro Met Ala Thr Gly Pro Ala Leu Ser 1345	1350	1355 1360
Ala Leu Glu Arg Leu Val Asn Thr Ser Ala Val Gln Arg Ser Val Thr 1365	1370	1375
Arg Met Asp Trp Ala Arg Phe Ala Pro Val Tyr Ala Ala Arg Gly Arg 1380	1385	1390
Arg Asn Leu Leu Ser Ala Leu Val Ala Glu Asp Glu Arg Ala Ala Ser 1395	1400	1405
Pro Pro Val Pro Thr Ala Asn Arg Ile Trp Arg Gly Leu Ser Val Ala 1410	1415	1420
Glu Ser Arg Ser Ala Leu Tyr Glu Leu Val Arg Gly Ile Val Ala Arg 1425	1430	1435 1440
Val Leu Gly Phe Ser Asp Pro Gly Ala Leu Asp Val Gly Arg Gly Phe 1445	1450	1455
Ala Glu Gln Gly Leu Asp Ser Leu Met Ala Leu Glu Ile Arg Asn Arg 1460	1465	1470
Leu Gln Arg Glu Leu Gly Glu Arg Leu Ser Ala Thr Leu Ala Phe Asp 1475	1480	1485
His Pro Thr Val Glu Arg Leu Val Ala His Leu Leu Thr Asp Val Leu 1490	1495	1500
Lys Leu Glu Asp Arg Ser Asp Thr Arg His Ile Arg Ser Val Ala Ala 1505	1510	1515 1520
Asp Asp Asp Ile Ala Ile Val Gly Ala Ala Cys Arg Phe Pro Gly Gly 1525	1530	1535
Asp Glu Gly Leu Glu Thr Tyr Trp Arg His Leu Ala Glu Gly Met Val 1540	1545	1550
Val Ser Thr Glu Val Pro Ala Asp Arg Trp Arg Ala Ala Asp Trp Tyr 1555	1560	1565
Asp Pro Asp Pro Glu Val Pro Gly Arg Thr Tyr Val Ala Lys Gly Ala 1570	1575	1580
Phe Leu Arg Asp Val Arg Ser Leu Asp Ala Ala Phe Phe Ala Ile Ser 1585	1590	1595 1600
Pro Arg Glu Ala Met Ser Leu Asp Pro Gln Gln Arg Leu Leu Leu Glu 1605	1610	1615
Val Ser Trp Glu Ala Ile Glu Arg Ala Gly Gln Asp Pro Met Ala Leu 1620	1625	1630
Arg Glu Ser Ala Thr Gly Val Phe Val Gly Met Ile Gly Ser Glu His 1635	1640	1645
Ala Glu Arg Val Gln Gly Leu Asp Asp Asp Ala Ala Leu Leu Tyr Gly 1650	1655	1660
Thr Thr Gly Asn Leu Leu Ser Val Ala Ala Gly Arg Leu Ser Phe Phe		

1665	1670	1675	1680
Leu Gly Leu His Gly Pro Thr Met Thr Val Asp Thr Ala Cys Ser Ser	1685	1690	1695
Ser Leu Val Ala Leu His Leu Ala Cys Gln Ser Leu Arg Leu Gly Glu	1700	1705	1710
Cys Asp Gln Ala Leu Ala Gly Gly Ser Ser Val Leu Leu Ser Pro Arg	1715	1720	1725
Ser Phe Val Ala Ala Ser Arg Met Arg Leu Leu Ser Pro Asp Gly Arg	1730	1735	1740
Cys Lys Thr Phe Ser Ala Ala Ala Asp Gly Phe Ala Arg Ala Glu Gly	1745	1750	1755
Cys Ala Val Val Val Leu Lys Arg Leu Arg Asp Ala Gln Arg Asp Arg	1765	1770	1775
Asp Pro Ile Leu Ala Val Val Arg Ser Thr Ala Ile Asn His Asp Gly	1780	1785	1790
Pro Ser Ser Gly Leu Thr Val Pro Ser Gly Pro Ala Gln Ala Leu	1795	1800	1805
Leu Arg Gln Ala Leu Ala Gln Ala Gly Val Ala Pro Ala Glu Val Asp	1810	1815	1820
Phe Val Glu Cys His Gly Thr Gly Thr Ala Leu Gly Asp Pro Ile Glu	1825	1830	1835
Val Gln Ala Leu Gly Ala Val Tyr Gly Arg Gly Arg Pro Ala Glu Arg	1845	1850	1855
Pro Leu Trp Leu Gly Ala Val Lys Ala Asn Leu Gly His Leu Glu Ala	1860	1865	1870
Ala Ala Gly Leu Ala Gly Val Leu Lys Val Leu Leu Ala Leu Glu His	1875	1880	1885
Glu Gln Ile Pro Ala Gln Pro Glu Leu Asp Glu Leu Asn Pro His Ile	1890	1895	1900
Pro Trp Ala Glu Leu Pro Val Ala Val Val Arg Arg Ala Val Pro Trp	1905	1910	1915
Pro Arg Gly Ala Arg Pro Arg Arg Ala Gly Val Ser Ala Phe Gly Leu	1925	1930	1935
Ser Gly Thr Asn Ala His Val Val Leu Glu Glu Ala Pro Ala Val Glu	1940	1945	1950
Pro Val Ala Ala Ala Pro Glu Arg Ala Ala Glu Leu Phe Val Leu Ser	1955	1960	1965
Ala Lys Ser Ala Ala Ala Leu Asp Ala Gln Ala Ala Arg Leu Arg Asp	1970	1975	1980
His Leu Glu Lys His Val Glu Leu Gly Leu Gly Asp Val Ala Phe Ser	1985	1990	1995
Leu Ala Thr Thr Arg Ser Ala Met Glu His Arg Leu Ala Val Ala Ala	2005	2010	2015
Ser Ser Arg Glu Ala Leu Arg Gly Ala Leu Ser Ala Ala Ala Gln Gly			

2020	2025	2030
His Thr Pro Pro Gly Ala Val Arg Gly Arg Ala Ser Gly Gly Ser Ala 2035 2040 2045		
Pro Lys Val Val Phe Val Phe Pro Gly Gln Gly Ser Gln Trp Val Gly 2050 2055 2060		
Met Gly Arg Lys Leu Met Ala Glu Glu Pro Val Phe Arg Ala Ala Leu 2065 2070 2075 2080		
Glu Gly Cys Asp Arg Ala Ile Glu Ala Glu Ala Gly Trp Ser Leu Leu 2085 2090 2095		
Gly Glu Leu Ser Ala Asp Glu Ala Ala Ser Gln Leu Gly Arg Ile Asp 2100 2105 2110		
Val Val Gln Pro Val Leu Phe Ala Met Glu Val Ala Leu Ser Ala Leu 2115 2120 2125		
Trp Arg Ser Trp Gly Val Glu Pro Glu Ala Val Val Gly His Ser Met 2130 2135 2140		
Gly Glu Val Ala Ala Ala His Val Ala Gly Ala Leu Ser Leu Glu Asp 2145 2150 2155 2160		
Ala Val Ala Ile Ile Cys Arg Arg Ser Arg Leu Leu Arg Arg Ile Ser 2165 2170 2175		
Gly Gln Gly Glu Met Ala Leu Val Glu Leu Ser Leu Glu Glu Ala Glu 2180 2185 2190		
Ala Ala Leu Arg Gly His Glu Gly Arg Leu Ser Val Ala Val Ser Asn 2195 2200 2205		
Ser Pro Arg Ser Thr Val Leu Ala Gly Glu Pro Ala Ala Leu Ser Glu 2210 2215 2220		
Val Leu Ala Ala Leu Thr Ala Lys Gly Val Phe Trp Arg Gln Val Lys 2225 2230 2235 2240		
Val Asp Val Ala Ser His Ser Pro Gln Val Asp Pro Leu Arg Glu Glu 2245 2250 2255		
Leu Ile Ala Ala Leu Gly Ala Ile Arg Pro Arg Ala Ala Ala Val Pro 2260 2265 2270		
Met Arg Ser Thr Val Thr Gly Gly Val Ile Ala Gly Pro Glu Leu Gly 2275 2280 2285		
Ala Ser Tyr Trp Ala Asp Asn Leu Arg Gln Pro Val Arg Phe Ala Ala 2290 2295 2300		
Ala Ala Gln Ala Leu Leu Glu Gly Gly Pro Ala Leu Phe Ile Glu Met 2305 2310 2315 2320		
Ser Pro His Pro Ile Leu Val Pro Pro Leu Asp Glu Ile Gln Thr Ala 2325 2330 2335		
Ala Glu Gln Gly Gly Ala Ala Val Gly Ser Leu Arg Arg Gly Gln Asp 2340 2345 2350		
Glu Arg Ala Thr Leu Leu Glu Ala Leu Gly Thr Leu Trp Ala Ser Gly 2355 2360 2365		
Tyr Pro Val Ser Trp Ala Arg Leu Phe Pro Ala Gly Gly Arg Arg Val		

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2725	2730	2735
Ser Val Val Tyr Arg Lys Val Ile Lys Arg Trp Met Glu His Leu Val 2740	2745	2750
Ala Ile Gly Ile Leu Val Gly Asp Gly Glu His Phe Val Ser Ser Gln 2755	2760	2765
Pro Leu Pro Glu Pro Asp Leu Ala Ala Val Leu Glu Glu Ala Gly Arg 2770	2775	2780
Val Phe Ala Asp Leu Pro Val Leu Phe Glu Trp Cys Lys Phe Ala Gly 2785	2790	2795
Glu Arg Leu Ala Asp Val Leu Thr Gly Lys Thr Leu Ala Leu Glu Ile 2805	2810	2815
Leu Phe Pro Gly Gly Ser Phe Asp Met Ala Glu Arg Ile Tyr Arg Asp 2820	2825	2830
Ser Pro Ile Ala Arg Tyr Ser Asn Gly Ile Val Arg Gly Val Val Glu 2835	2840	2845
Ser Ala Ala Arg Val Val Ala Pro Ser Gly Met Phe Ser Ile Leu Glu 2850	2855	2860
Ile Gly Ala Gly Thr Gly Ala Thr Thr Ala Ala Val Leu Pro Val Leu 2865	2870	2875
Leu Pro Asp Arg Thr Glu Tyr His Phe Thr Asp Val Ser Pro Leu Phe 2885	2890	2895
Leu Ala Arg Ala Glu Gln Arg Phe Arg Asp Tyr Pro Phe Leu Lys Tyr 2900	2905	2910
Gly Ile Leu Asp Val Asp Gln Glu Pro Ala Gly Gln Gly Tyr Ala His 2915	2920	2925
Gln Arg Phe Asp Val Ile Val Ala Ala Asn Val Ile His Ala Thr Arg 2930	2935	2940
Asp Ile Arg Ala Thr Ala Lys Arg Leu Leu Ser Leu Leu Ala Pro Gly 2945	2950	2955
Gly Leu Leu Val Leu Val Glu Gly Thr Gly His Pro Ile Trp Phe Asp 2965	2970	2975
Ile Thr Thr Gly Leu Ile Glu Gly Trp Gln Lys Tyr Glu Asp Asp Leu 2980	2985	2990
Arg Ile Asp His Pro Leu Leu Pro Ala Arg Thr Trp Cys Asp Val Leu 2995	3000	3005
Arg Arg Val Gly Phe Ala Asp Ala Val Ser Leu Pro Gly Asp Gly Ser 3010	3015	3020
Pro Ala Gly Ile Leu Gly Gln His Val Ile Leu Ser Arg Ala Pro Gly 3025	3030	3035
Ile Ala Gly Ala Ala Cys Asp Ser Ser Gly Glu Ser Ala Thr Glu Ser 3045	3050	3055
Pro Ala Ala Arg Ala Val Arg Gln Glu Trp Ala Asp Gly Ser Ala Asp 3060	3065	3070
Val Val His Arg Met Ala Leu Glu Arg Met Tyr Phe His Arg Arg Pro		

3075	3080	3085
Gly Arg Gln Val Trp Val His Gly Arg Leu Arg Thr Gly Gly Gly Ala 3090 3095 3100		
Phe Thr Lys Ala Leu Ala Gly Asp Leu Leu Leu Phe Glu Asp Thr Gly 3105 3110 3115 3120		
Gln Val Val Ala Glu Val Gln Gly Leu Arg Leu Pro Gln Leu Glu Ala 3125 3130 3135		
Ser Ala Phe Ala Pro Arg Asp Pro Arg Glu Glu Trp Leu Tyr Ala Leu 3140 3145 3150		
Glu Trp Gln Arg Lys Asp Pro Ile Pro Glu Ala Pro Ala Ala Ala Ser 3155 3160 3165		
Ser Ser Ser Ala Gly Ala Trp Leu Val Leu Met Asp Gln Gly Gly Thr 3170 3175 3180		
Gly Ala Ala Leu Val Ser Leu Leu Glu Gly Arg Gly Glu Ala Cys Val 3185 3190 3195 3200		
Arg Val Ile Ala Gly Thr Ala Tyr Ala Cys Leu Ala Pro Gly Leu Tyr 3205 3210 3215		
Gln Val Asp Pro Ala Gln Pro Asp Gly Phe His Thr Leu Leu Arg Asp 3220 3225 3230		
Ala Phe Gly Glu Asp Arg Ile Cys Arg Ala Val Val His Met Trp Ser 3235 3240 3245		
Leu Asp Ala Thr Ala Ala Gly Glu Arg Ala Thr Ala Glu Ser Leu Gln 3250 3255 3260		
Ala Asp Gln Leu Leu Gly Ser Leu Ser Ala Leu Ser Leu Val Gln Ala 3265 3270 3275 3280		
Leu Val Arg Arg Arg Trp Arg Asn Met Pro Arg Leu Trp Leu Leu Thr 3285 3290 3295		
Arg Ala Val His Ala Val Gly Ala Glu Asp Ala Ala Ala Ser Val Ala 3300 3305 3310		
Gln Ala Pro Val Trp Gly Leu Gly Arg Thr Leu Ala Leu Glu His Pro 3315 3320 3325		
Glu Leu Arg Cys Thr Leu Val Asp Val Asn Pro Ala Pro Ser Pro Glu 3330 3335 3340		
Asp Ala Ala Ala Leu Ala Val Glu Leu Gly Ala Ser Asp Arg Glu Asp 3345 3350 3355 3360		
Gln Val Ala Leu Arg Ser Asp Gly Arg Tyr Val Ala Arg Leu Val Arg 3365 3370 3375		
Ser Ser Phe Ser Gly Lys Pro Ala Thr Asp Cys Gly Ile Arg Ala Asp 3380 3385 3390		
Gly Ser Tyr Val Ile Thr Asp Gly Met Gly Arg Val Gly Leu Ser Val 3395 3400 3405		
Ala Gln Trp Met Val Met Gln Gly Ala Arg His Val Val Leu Val Asp 3410 3415 3420		
Arg Gly Gly Ala Ser Glu Ala Ser Arg Asp Ala Leu Arg Ser Met Ala		

3425	3430	3435	3440
Glu Ala Gly	Glu Val Gln Ile Val	Glu Ala Asp Val Ala Arg Arg	
3445	3450	3455	
Asp Asp Val Ala Arg	Leu Leu Ser Lys Ile Glu Pro Ser Met Pro Pro		
3460	3465	3470	
Leu Arg Gly Ile Val Tyr Val Asp Gly Thr Phe Gln Gly Asp Ser Ser			
3475	3480	3485	
Met Leu Glu Leu Asp Ala Arg Arg Phe Lys Glu Trp Met Tyr Pro Lys			
3490	3495	3500	
Val Leu Gly Ala Trp Asn Leu His Ala Leu Thr Arg Asp Arg Ser Leu			
3505	3510	3515	3520
Asp Phe Phe Val Leu Tyr Ser Ser Gly Thr Ser Leu Leu Gly Leu Pro			
3525	3530	3535	
Gly Gln Gly Ser Arg Ala Ala Gly Asp Ala Phe Leu Asp Ala Ile Ala			
3540	3545	3550	
His His Arg Cys Lys Val Gly Leu Thr Ala Met Ser Ile Asn Trp Gly			
3555	3560	3565	
Leu Leu Ser Glu Ala Ser Ser Pro Ala Thr Pro Asn Asp Gly Gly Ala			
3570	3575	3580	
Arg Leu Glu Tyr Arg Gly Met Glu Gly Leu Thr Leu Glu Gln Gly Ala			
3585	3590	3595	3600
Ala Ala Leu Gly Arg Leu Leu Ala Arg Pro Arg Ala Gln Val Gly Val			
3605	3610	3615	
Met Arg Leu Asn Leu Arg Gln Trp Leu Glu Phe Tyr Pro Asn Ala Ala			
3620	3625	3630	
Arg Leu Ala Leu Trp Ala Glu Leu Leu Lys Glu Arg Asp Arg Ala Asp			
3635	3640	3645	
Arg Gly Ala Ser Asn Ala Ser Asn Leu Arg Glu Ala Leu Gln Ser Ala			
3650	3655	3660	
Arg Pro Glu Asp Arg Gln Leu Ile Leu Glu Lys His Leu Ser Glu Leu			
3665	3670	3675	3680
Leu Gly Arg Gly Leu Arg Leu Pro Pro Glu Arg Ile Glu Arg His Val			
3685	3690	3695	
Pro Phe Ser Asn Leu Gly Met Asp Ser Leu Ile Gly Leu Glu Leu Arg			
3700	3705	3710	
Asn Arg Ile Glu Ala Ala Leu Gly Ile Thr Val Pro Ala Thr Leu Leu			
3715	3720	3725	
Trp Thr Tyr Pro Asn Val Ala Ala Leu Ser Gly Ser Leu Leu Asp Ile			
3730	3735	3740	
Leu Phe Pro Asn Ala Gly Ala Thr His Ala Pro Ala Thr Glu Arg Glu			
3745	3750	3755	3760
Lys Ser Phe Glu Asn Asp Ala Ala Asp Leu Glu Ala Leu Arg Gly Met			
3765	3770	3775	
Thr Asp Glu Gln Lys Asp Ala Leu Leu Ala Glu Lys Leu Ala Gln Leu			

3780

3785

3790

Ala Gln Ile Val Gly Glu
3795

<210> 7

<211> 2439

<212> PRT

<213> Sorangium cellulosum

<400> 7

Met Ala Thr Thr Asn Ala Gly Lys Leu Glu His Ala Leu Leu Leu Met
1 5 10 15

Asp Lys Leu Ala Lys Lys Asn Ala Ser Leu Glu Gln Glu Arg Thr Glu
20 25 30

Pro Ile Ala Ile Val Gly Ile Gly Cys Arg Phe Pro Gly Gly Ala Asp
35 40 45

Thr Pro Glu Ala Phe Trp Glu Leu Leu Asp Ser Gly Arg Asp Ala Val
50 55 60

Gln Pro Leu Asp Arg Arg Trp Ala Leu Val Gly Val His Pro Ser Glu
65 70 75 80

Glu Val Pro Arg Trp Ala Gly Leu Leu Thr Glu Ala Val Asp Gly Phe
85 90 95

Asp Ala Ala Phe Phe Gly Thr Ser Pro Arg Glu Ala Arg Ser Leu Asp
100 105 110

Pro Gln Gln Arg Leu Leu Leu Glu Val Thr Trp Glu Gly Leu Glu Asp
115 120 125

Ala Gly Ile Ala Pro Gln Ser Leu Asp Gly Ser Arg Thr Gly Val Phe
130 135 140

Leu Gly Ala Cys Ser Ser Asp Tyr Ser His Thr Val Ala Gln Gln Arg
145 150 155 160

Arg Glu Glu Gln Asp Ala Tyr Asp Ile Thr Gly Asn Thr Leu Ser Val
165 170 175

Ala Ala Gly Arg Leu Ser Tyr Thr Leu Gly Leu Gln Gly Pro Cys Leu
180 185 190

Thr Val Asp Thr Ala Cys Ser Ser Ser Leu Val Ala Ile His Leu Ala
195 200 205

Cys Arg Ser Leu Arg Ala Arg Glu Ser Asp Leu Ala Leu Ala Gly Gly
210 215 220

Val Asn Met Leu Leu Ser Ser Lys Thr Met Ile Met Leu Gly Arg Ile
225 230 235 240

Gln Ala Leu Ser Pro Asp Gly His Cys Arg Thr Phe Asp Ala Ser Ala
245 250 255

Asn Gly Phe Val Arg Gly Glu Gly Cys Gly Met Val Val Leu Lys Arg
260 265 270

Leu Ser Asp Ala Gln Arg His Gly Asp Arg Ile Trp Ala Leu Ile Arg
275 280 285

Gly Ser Ala Met Asn Gln Asp Gly Arg Ser Thr Gly Leu Met Ala Pro
 290 295 300
 Asn Val Leu Ala Gln Glu Ala Leu Leu Arg Glu Ala Leu Gln Ser Ala
 305 310 315 320
 Arg Val Asp Ala Gly Ala Ile Gly Tyr Val Glu Thr His Gly Thr Gly
 325 330 335
 Thr Ser Leu Gly Asp Pro Ile Glu Val Glu Ala Leu Arg Ala Val Leu
 340 345 350
 Gly Pro Ala Arg Ala Asp Gly Ser Arg Cys Val Leu Gly Ala Val Lys
 355 360 365
 Thr Asn Leu Gly His Leu Glu Gly Ala Ala Gly Val Ala Gly Leu Ile
 370 375 380
 Lys Ala Ala Leu Ala Leu His His Glu Leu Ile Pro Arg Asn Leu His
 385 390 395 400
 Phe His Thr Leu Asn Pro Arg Ile Arg Ile Glu Gly Thr Ala Leu Ala
 405 410 415
 Leu Ala Thr Glu Pro Val Pro Trp Pro Arg Ala Gly Arg Pro Arg Phe
 420 425 430
 Ala Gly Val Ser Ala Phe Gly Leu Ser Gly Thr Asn Val His Val Val
 435 440 445
 Leu Glu Glu Ala Pro Ala Thr Val Leu Ala Pro Ala Thr Pro Gly Arg
 450 455 460
 Ser Ala Glu Leu Leu Val Leu Ser Ala Lys Ser Ala Ala Ala Leu Asp
 465 470 475 480
 Ala Gln Ala Ala Arg Leu Ser Ala His Ile Ala Ala Tyr Pro Glu Gln
 485 490 495
 Gly Leu Gly Asp Val Ala Phe Ser Leu Val Ser Thr Arg Ser Pro Met
 500 505 510
 Glu His Arg Leu Ala Val Ala Ala Thr Ser Arg Glu Ala Leu Arg Ser
 515 520 525
 Ala Leu Glu Val Ala Ala Gln Gly Gln Thr Pro Ala Gly Ala Ala Arg
 530 535 540
 Gly Arg Ala Ala Ser Ser Pro Gly Lys Leu Ala Phe Leu Phe Ala Gly
 545 550 555 560
 Gln Gly Ala Gln Val Pro Gly Met Gly Arg Gly Leu Trp Glu Ala Trp
 565 570 575
 Pro Ala Phe Arg Glu Thr Phe Asp Arg Cys Val Thr Leu Phe Asp Arg
 580 585 590
 Glu Leu His Gln Pro Leu Cys Glu Val Met Trp Ala Glu Pro Gly Ser
 595 600 605
 Ser Arg Ser Ser Leu Leu Asp Gln Thr Ala Phe Thr Gln Pro Ala Leu
 610 615 620
 Phe Ala Leu Glu Tyr Ala Leu Ala Ala Leu Phe Arg Ser Trp Gly Val
 625 630 635 640

Glu Pro Glu Leu Val Ala Gly His Ser Leu Gly Glu Leu Val Ala Ala
 645 650 655
 Cys Val Ala Gly Val Phe Ser Leu Glu Asp Ala Val Arg Leu Val Val
 660 665 670
 Ala Arg Gly Arg Leu Met Gln Ala Leu Pro Ala Gly Gly Ala Met Val
 675 680 685
 Ser Ile Ala Ala Pro Glu Ala Asp Val Ala Ala Ala Val Ala Pro His
 690 695 700
 Ala Ala Leu Val Ser Ile Ala Ala Val Asn Gly Pro Glu Gln Val Val
 705 710 715 720
 Ile Ala Gly Ala Glu Lys Phe Val Gln Gln Ile Ala Ala Ala Phe Ala
 725 730 735
 Ala Arg Gly Ala Arg Thr Lys Pro Leu His Val Ser His Ala Phe His
 740 745 750
 Ser Pro Leu Met Asp Pro Met Leu Glu Ala Phe Arg Arg Val Thr Glu
 755 760 765
 Ser Val Thr Tyr Arg Arg Pro Ser Ile Ala Leu Val Ser Asn Leu Ser
 770 775 780
 Gly Lys Pro Cys Thr Asp Glu Val Ser Ala Pro Gly Tyr Trp Val Arg
 785 790 795 800
 His Ala Arg Glu Ala Val Arg Phe Ala Asp Gly Val Lys Ala Leu His
 805 810 815
 Ala Ala Gly Ala Gly Leu Phe Val Glu Val Gly Pro Lys Pro Thr Leu
 820 825 830
 Leu Gly Leu Val Pro Ala Cys Leu Pro Asp Ala Arg Pro Val Leu Leu
 835 840 845
 Pro Ala Ser Arg Ala Gly Arg Asp Glu Ala Ala Ser Ala Leu Glu Ala
 850 855 860
 Leu Gly Gly Phe Trp Val Val Gly Gly Ser Val Thr Trp Ser Gly Val
 865 870 875 880
 Phe Pro Ser Gly Gly Arg Arg Val Pro Leu Pro Thr Tyr Pro Trp Gln
 885 890 895
 Arg Glu Arg Tyr Trp Ile Glu Ala Pro Val Asp Arg Glu Ala Asp Gly
 900 905 910
 Thr Gly Arg Ala Arg Ala Gly Gly His Pro Leu Leu Gly Glu Val Phe
 915 920 925
 Ser Val Ser Thr His Ala Gly Leu Arg Leu Trp Glu Thr Thr Leu Asp
 930 935 940
 Arg Lys Arg Leu Pro Trp Leu Gly Glu His Arg Ala Gln Gly Glu Val
 945 950 955 960
 Val Phe Pro Gly Ala Gly Tyr Leu Glu Met Ala Leu Ser Ser Gly Ala
 965 970 975
 Glu Ile Leu Gly Asp Gly Pro Ile Gln Val Thr Asp Val Val Leu Ile
 980 985 990

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Glu Thr Leu Thr Phe Ala Gly Asp Thr Ala Val Pro Val Gln Val Val
 995 1000 1005
 Thr Thr Glu Glu Arg Pro Gly Arg Leu Arg Phe Gln Val Ala Ser Arg
 1010 1015 1020
 Glu Pro Gly Glu Arg Arg Ala Pro Phe Arg Ile His Ala Arg Gly Val
 1025 1030 1035 1040
 Leu Arg Arg Ile Gly Arg Val Glu Thr Pro Ala Arg Ser Asn Leu Ala
 1045 1050 1055
 Ala Leu Arg Ala Arg Leu His Ala Ala Val Pro Ala Ala Ala Ile Tyr
 1060 1065 1070
 Gly Ala Leu Ala Glu Met Gly Leu Gln Tyr Gly Pro Ala Leu Arg Gly
 1075 1080 1085
 Leu Ala Glu Leu Trp Arg Gly Glu Gly Glu Ala Leu Gly Arg Val Arg
 1090 1095 1100
 Leu Pro Glu Ala Ala Gly Ser Ala Thr Ala Tyr Gln Leu His Pro Val
 1105 1110 1115 1120
 Leu Leu Asp Ala Cys Val Gln Met Ile Val Gly Ala Phe Ala Asp Arg
 1125 1130 1135
 Asp Glu Ala Thr Pro Trp Ala Pro Val Glu Val Gly Ser Val Arg Leu
 1140 1145 1150
 Phe Gln Arg Ser Pro Gly Glu Leu Trp Cys His Ala Arg Val Val Ser
 1155 1160 1165
 Asp Gly Gln Gln Ala Ser Ser Arg Trp Ser Ala Asp Phe Glu Leu Met
 1170 1175 1180
 Asp Gly Thr Gly Ala Val Val Ala Glu Ile Ser Arg Leu Val Val Glu
 1185 1190 1195 1200
 Arg Leu Ala Ser Gly Val Arg Arg Arg Asp Ala Asp Asp Trp Phe Leu
 1205 1210 1215
 Glu Leu Asp Trp Glu Pro Ala Ala Leu Gly Gly Pro Lys Ile Thr Ala
 1220 1225 1230
 Gly Arg Trp Leu Leu Leu Gly Glu Gly Gly Gly Leu Gly Arg Ser Leu
 1235 1240 1245
 Cys Ser Ala Leu Lys Ala Ala Gly His Val Val Val His Ala Ala Gly
 1250 1255 1260
 Asp Asp Thr Ser Thr Ala Gly Met Arg Ala Leu Leu Ala Asn Ala Phe
 1265 1270 1275 1280
 Asp Gly Gln Ala Pro Thr Ala Val Val His Leu Ser Ser Leu Asp Gly
 1285 1290 1295
 Gly Gly Gln Leu Gly Pro Gly Leu Gly Ala Gln Gly Ala Leu Asp Ala
 1300 1305 1310
 Pro Arg Ser Pro Asp Val Asp Ala Asp Ala Leu Glu Ser Ala Leu Met
 1315 1320 1325
 Arg Gly Cys Asp Ser Val Leu Ser Leu Val Gln Ala Leu Val Gly Met
 1330 1335 1340

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Asp Leu Arg Asn Ala Pro Arg Leu Trp Leu Leu Thr Arg Gly Ala Gln
 1345 1350 1355 1360
 Ala Ala Ala Ala Gly Asp Val Ser Val Val Gln Ala Pro Leu Leu Gly
 1365 1370 1375
 Leu Gly Arg Thr Ile Ala Leu Glu His Ala Glu Leu Arg Cys Ile Ser
 1380 1385 1390
 Val Asp Leu Asp Pro Ala Glu Pro Glu Gly Glu Ala Asp Ala Leu Leu
 1395 1400 1405
 Ala Glu Leu Leu Ala Asp Asp Ala Glu Glu Glu Val Ala Leu Arg Gly
 1410 1415 1420
 Gly Asp Arg Leu Val Ala Arg Leu Val His Arg Leu Pro Asp Ala Gln
 1425 1430 1435 1440
 Arg Arg Glu Lys Val Glu Pro Ala Gly Asp Arg Pro Phe Arg Leu Glu
 1445 1450 1455
 Ile Asp Glu Pro Gly Ala Leu Asp Gln Leu Val Leu Arg Ala Thr Gly
 1460 1465 1470
 Arg Arg Ala Pro Gly Pro Gly Glu Val Glu Ile Ser Val Glu Ala Ala
 1475 1480 1485
 Gly Leu Asp Ser Ile Asp Ile Gln Leu Ala Leu Gly Val Ala Pro Asn
 1490 1495 1500
 Asp Leu Pro Gly Glu Glu Ile Glu Pro Leu Val Leu Gly Ser Glu Cys
 1505 1510 1515 1520
 Ala Gly Arg Ile Val Ala Val Gly Glu Gly Val Asn Gly Leu Val Val
 1525 1530 1535
 Gly Gln Pro Val Ile Ala Leu Ala Ala Gly Val Phe Ala Thr His Val
 1540 1545 1550
 Thr Thr Ser Ala Thr Leu Val Leu Pro Arg Pro Leu Gly Leu Ser Ala
 1555 1560 1565
 Thr Glu Ala Ala Ala Met Pro Leu Ala Tyr Leu Thr Ala Trp Tyr Ala
 1570 1575 1580
 Leu Asp Lys Val Ala His Leu Gln Ala Gly Glu Arg Val Leu Ile His
 1585 1590 1595 1600
 Ala Glu Ala Gly Gly Val Gly Leu Cys Ala Val Arg Trp Ala Gln Arg
 1605 1610 1615
 Val Gly Ala Glu Val Tyr Ala Thr Ala Asp Thr Pro Glu Asn Arg Ala
 1620 1625 1630
 Tyr Leu Glu Ser Leu Gly Val Arg Tyr Val Ser Asp Ser Arg Ser Gly
 1635 1640 1645
 Arg Phe Val Thr Asp Val His Ala Trp Thr Asp Gly Glu Gly Val Asp
 1650 1655 1660
 Val Val Leu Asp Ser Leu Ser Gly Glu Arg Ile Asp Lys Ser Leu Met
 1665 1670 1675 1680
 Val Leu Arg Ala Cys Gly Arg Leu Val Lys Leu Gly Arg Arg Asp Asp
 1685 1690 1695

Cys Ala Asp Thr Gln Pro Gly Leu Pro Pro Leu Leu Arg Asn Phe Ser
 1700 1705 1710
 Phe Ser Gln Val Asp Leu Arg Gly Met Met Leu Asp Gln Pro Ala Arg
 1715 1720 1725
 Ile Arg Ala Leu Leu Asp Glu Leu Phe Gly Leu Val Ala Ala Gly Ala
 1730 1735 1740
 Ile Ser Pro Leu Gly Ser Gly Leu Arg Val Gly Gly Ser Leu Thr Pro
 1745 1750 1755 1760
 Pro Pro Val Glu Thr Phe Pro Ile Ser Arg Ala Ala Glu Ala Phe Arg
 1765 1770 1775
 Arg Met Ala Gln Gly Gln His Leu Gly Lys Leu Val Leu Thr Leu Asp
 1780 1785 1790
 Asp Pro Glu Val Arg Ile Arg Ala Pro Ala Glu Ser Ser Val Ala Val
 1795 1800 1805
 Arg Ala Asp Gly Thr Tyr Leu Val Thr Gly Gly Leu Gly Gly Leu Gly
 1810 1815 1820
 Leu Arg Val Ala Gly Trp Leu Ala Glu Arg Gly Ala Gly Gln Leu Val
 1825 1830 1835 1840
 Leu Val Gly Arg Ser Gly Ala Ala Ser Ala Glu Gln Arg Ala Ala Val
 1845 1850 1855
 Ala Ala Leu Glu Ala His Gly Ala Arg Val Thr Val Ala Lys Ala Asp
 1860 1865 1870
 Val Ala Asp Arg Ser Gln Ile Glu Arg Val Leu Arg Glu Val Thr Ala
 1875 1880 1885
 Ser Gly Met Pro Leu Arg Gly Val Val His Ala Ala Gly Leu Val Asp
 1890 1895 1900
 Asp Gly Leu Leu Met Gln Gln Thr Pro Ala Arg Phe Arg Thr Val Met
 1905 1910 1915 1920
 Gly Pro Lys Val Gln Gly Ala Leu His Leu His Thr Leu Thr Arg Glu
 1925 1930 1935
 Ala Pro Leu Ser Phe Phe Val Leu Tyr Ala Ser Ala Ala Gly Leu Phe
 1940 1945 1950
 Gly Ser Pro Gly Gln Gly Asn Tyr Ala Ala Ala Asn Ala Phe Leu Asp
 1955 1960 1965
 Ala Leu Ser His His Arg Arg Ala Gln Gly Leu Pro Ala Leu Ser Ile
 1970 1975 1980
 Asp Trp Gly Met Phe Thr Glu Val Gly Met Ala Val Ala Gln Glu Asn
 1985 1990 1995 2000
 Arg Gly Ala Arg Gln Ile Ser Arg Gly Met Arg Gly Ile Thr Pro Asp
 2005 2010 2015
 Glu Gly Leu Ser Ala Leu Ala Arg Leu Leu Glu Gly Asp Arg Val Gln
 2020 2025 2030
 Thr Gly Val Ile Pro Ile Thr Pro Arg Gln Trp Val Glu Phe Tyr Pro
 2035 2040 2045

Ala Thr Ala Ala Ser Arg Arg Leu Ser Arg Leu Val Thr Thr Gln Arg
 2050 2055 2060
 Ala Val Ala Asp Arg Thr Ala Gly Asp Arg Asp Leu Leu Glu Gln Leu
 2065 2070 2075 2080
 Ala Ser Ala Glu Pro Ser Ala Arg Ala Gly Leu Leu Gln Asp Val Val
 2085 2090 2095
 Arg Val Gln Val Ser His Val Leu Arg Leu Pro Glu Asp Lys Ile Glu
 2100 2105 2110
 Val Asp Ala Pro Leu Ser Ser Met Gly Met Asp Ser Leu Met Ser Leu
 2115 2120 2125
 Glu Leu Arg Asn Arg Ile Glu Ala Ala Leu Gly Val Ala Ala Pro Ala
 2130 2135 2140
 Ala Leu Gly Trp Thr Tyr Pro Thr Val Ala Ala Ile Thr Arg Trp Leu
 2145 2150 2155 2160
 Leu Asp Asp Ala Leu Val Val Arg Leu Gly Gly Gly Ser Asp Thr Asp
 2165 2170 2175
 Glu Ser Thr Ala Ser Ala Gly Ser Phe Val His Val Leu Arg Phe Arg
 2180 2185 2190
 Pro Val Val Lys Pro Arg Ala Arg Leu Phe Cys Phe His Gly Ser Gly
 2195 2200 2205
 Gly Ser Pro Glu Gly Phe Arg Ser Trp Ser Glu Lys Ser Glu Trp Ser
 2210 2215 2220
 Asp Leu Glu Ile Val Ala Met Trp His Asp Arg Ser Leu Ala Ser Glu
 2225 2230 2235 2240
 Asp Ala Pro Gly Lys Lys Tyr Val Gln Glu Ala Ala Ser Leu Ile Gln
 2245 2250 2255
 His Tyr Ala Asp Ala Pro Phe Ala Leu Val Gly Phe Ser Leu Gly Val
 2260 2265 2270
 Arg Phe Val Met Gly Thr Ala Val Glu Leu Ala Ser Arg Ser Gly Ala
 2275 2280 2285
 Pro Ala Pro Leu Ala Val Phe Thr Leu Gly Gly Ser Leu Ile Ser Ser
 2290 2295 2300
 Ser Glu Ile Thr Pro Glu Met Glu Thr Asp Ile Ile Ala Lys Leu Phe
 2305 2310 2315 2320
 Phe Arg Asn Ala Ala Gly Phe Val Arg Ser Thr Gln Gln Val Gln Ala
 2325 2330 2335
 Asp Ala Arg Ala Asp Lys Val Ile Thr Asp Thr Met Val Ala Pro Ala
 2340 2345 2350
 Pro Gly Asp Ser Lys Glu Pro Pro Val Lys Ile Ala Val Pro Ile Val
 2355 2360 2365
 Ala Ile Ala Gly Ser Asp Asp Val Ile Val Pro Pro Ser Asp Val Gln
 2370 2375 2380
 Asp Leu Gln Ser Arg Thr Thr Glu Arg Phe Tyr Met His Leu Leu Pro
 2385 2390 2395 2400

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Gly Asp His Glu Phe Leu Val Asp Arg Gly Arg Glu Ile Met His Ile
 2405 2410 2415

Val Asp Ser His Leu Asn Pro Leu Leu Ala Ala Arg Thr Thr Ser Ser
 2420 2425 2430

Gly Pro Ala Phe Glu Ala Lys
 2435

<210> 8

<211> 419

<212> PRT

<213> Sorangium cellulosum

<400> 8

Met Thr Gln Glu Gln Ala Asn Gln Ser Glu Thr Lys Pro Ala Phe Asp
 1 5 10 15

Phe Lys Pro Phe Ala Pro Gly Tyr Ala Glu Asp Pro Phe Pro Ala Ile
 20 25 30

Glu Arg Leu Arg Glu Ala Thr Pro Ile Phe Tyr Trp Asp Glu Gly Arg
 35 40 45

Ser Trp Val Leu Thr Arg Tyr His Asp Val Ser Ala Val Phe Arg Asp
 50 55 60

Glu Arg Phe Ala Val Ser Arg Glu Glu Trp Glu Ser Ser Ala Glu Tyr
 65 70 75 80

Ser Ser Ala Ile Pro Glu Leu Ser Asp Met Lys Lys Tyr Gly Leu Phe
 85 90 95

Gly Leu Pro Pro Glu Asp His Ala Arg Val Arg Lys Leu Val Asn Pro
 100 105 110

Ser Phe Thr Ser Arg Ala Ile Asp Leu Leu Arg Ala Glu Ile Gln Arg
 115 120 125

Thr Val Asp Gln Leu Leu Asp Ala Arg Ser Gly Gln Glu Glu Phe Asp
 130 135 140

Val Val Arg Asp Tyr Ala Glu Gly Ile Pro Met Arg Ala Ile Ser Ala
 145 150 155 160

Leu Leu Lys Val Pro Ala Glu Cys Asp Glu Lys Phe Arg Arg Phe Gly
 165 170 175

Ser Ala Thr Ala Arg Ala Leu Gly Val Gly Leu Val Pro Gln Val Asp
 180 185 190

Glu Glu Thr Lys Thr Leu Val Ala Ser Val Thr Glu Gly Leu Ala Leu
 195 200 205

Leu His Asp Val Leu Asp Glu Arg Arg Arg Asn Pro Leu Glu Asn Asp
 210 215 220

Val Leu Thr Met Leu Leu Gln Ala Glu Ala Asp Gly Ser Arg Leu Ser
 225 230 235 240

Thr Lys Glu Leu Val Ala Leu Val Gly Ala Ile Ile Ala Ala Gly Thr
 245 250 255

Asp Thr Thr Ile Tyr Leu Ile Ala Phe Ala Val Leu Asn Leu Leu Arg
 260 265 270

Ser Pro Glu Ala Leu Glu Leu Val Lys Ala Glu Pro Gly Leu Met Arg
275 280 285

Asn Ala Leu Asp Glu Val Leu Arg Phe Asp Asn Ile Leu Arg Ile Gly
290 295 300

Thr Val Arg Phe Ala Arg Gln Asp Leu Glu Tyr Cys Gly Ala Ser Ile
305 310 315 320

Lys Lys Gly Glu Met Val Phe Leu Leu Ile Pro Ser Ala Leu Arg Asp
325 330 335

Gly Thr Val Phe Ser Arg Pro Asp Val Phe Asp Val Arg Arg Asp Thr
340 345 350

Gly Ala Ser Leu Ala Tyr Gly Arg Gly Pro His Val Cys Pro Gly Val
355 360 365

Ser Leu Ala Arg Leu Glu Ala Glu Ile Ala Val Gly Thr Ile Phe Arg
370 375 380

Arg Phe Pro Glu Met Lys Leu Lys Glu Thr Pro Val Phe Gly Tyr His
385 390 395 400

Pro Ala Phe Arg Asn Ile Glu Ser Leu Asn Val Ile Leu Lys Pro Ser
405 410 415

Lys Ala Gly

<210> 9

<211> 607

<212> PRT

<213> Sorangium cellulosum

<400> 9

Ala Ser Leu Asp Ala Leu Phe Ala Arg Ala Thr Ser Ala Arg Val Leu
1 5 10 15

Asp Asp Gly His Gly Arg Ala Thr Glu Arg His Val Leu Ala Glu Ala
20 25 30

Arg Gly Ile Glu Asp Leu Arg Ala Leu Arg Glu His Leu Arg Ile Gln
35 40 45

Glu Gly Gly Pro Ser Phe His Cys Met Cys Leu Gly Asp Leu Thr Val
50 55 60

Glu Leu Leu Ala His Asp Gln Pro Leu Ala Ser Ile Ser Phe His His
65 70 75 80

Ala Arg Ser Leu Arg His Pro Asp Trp Thr Ser Asp Ala Met Leu Val
85 90 95

Asp Gly Pro Ala Leu Val Arg Trp Leu Ala Ala Arg Gly Ala Pro Gly
100 105 110

Pro Leu Arg Glu Tyr Glu Glu Glu Arg Glu Arg Ala Arg Thr Ala Gln
115 120 125

Glu Ala Arg Arg Leu Trp Leu Ala Ala Ala Pro Pro Cys Phe Ala Pro
130 135 140

Asp Leu Pro Arg Phe Glu Asp Asp Ala Asn Gly Leu Pro Leu Gly Pro

145 150 155 160
 Met Ser Pro Glu Val Ala Glu Ala Glu Arg Arg Leu Arg Ala Ser Tyr
 165 170 175
 Ala Thr Pro Glu Leu Ala Cys Ala Ala Leu Leu Ala Trp Leu Gly Thr
 180 185 190
 Gly Ala Gly Pro Trp Ser Gly Tyr Pro Ala Tyr Glu Met Leu Pro Glu
 195 200 205
 Asn Leu Leu Leu Gly Phe Gly Leu Pro Thr Ala Ile Ala Ala Ala Ser
 210 215 220
 Ala Pro Gly Thr Ser Glu Ala Ala Leu Arg Gly Ala Ala Arg Leu Phe
 225 230 235 240
 Ala Ser Trp Glu Val Val Ser Ser Lys Lys Ser Gln Leu Gly Asn Ile
 245 250 255
 Pro Glu Ala Leu Trp Glu Arg Leu Arg Thr Ile Val Arg Ala Met Gly
 260 265 270
 Asn Ala Asp Asn Leu Ser Arg Phe Glu Arg Ala Glu Ala Ile Ala Ala
 275 280 285
 Glu Val Arg Arg Leu Arg Ala Gln Pro Ala Pro Phe Ala Ala Gly Ala
 290 295 300
 Gly Leu Ala Val Ala Gly Val Ser Ser Ser Gly Arg Leu Ser Gly Leu
 305 310 315 320
 Val Thr Asp Gly Asp Ala Leu Tyr Ser Gly Asp Gly Asn Asp Ile Val
 325 330 335
 Met Phe Gln Pro Gly Arg Ile Ser Pro Val Val Leu Leu Ala Gly Thr
 340 345 350
 Asp Pro Phe Phe Glu Leu Ala Pro Pro Leu Ser Gln Met Leu Phe Val
 355 360 365
 Ala His Ala Asn Ala Gly Thr Ile Ser Lys Val Leu Thr Glu Gly Ser
 370 375 380
 Pro Leu Ile Val Met Ala Arg Asn Gln Ala Arg Pro Met Ser Leu Val
 385 390 395 400
 His Ala Arg Gly Phe Met Ala Trp Val Asn Gln Ala Met Val Pro Asp
 405 410 415
 Pro Glu Arg Gly Ala Pro Phe Val Val Gln Arg Ser Thr Ile Met Glu
 420 425 430
 Phe Glu His Pro Thr Pro Arg Cys Leu His Glu Pro Ala Gly Ser Ala
 435 440 445
 Phe Ser Leu Ala Cys Asp Glu Glu His Leu Tyr Trp Cys Glu Leu Ser
 450 455 460
 Ala Gly Arg Leu Glu Leu Trp Arg His Pro His His Arg Pro Gly Ala
 465 470 475 480
 Pro Ser Arg Phe Ala Tyr Leu Gly Glu His Pro Ile Ala Ala Thr Trp
 485 490 495
 Tyr Pro Ser Leu Thr Leu Asn Ala Thr His Val Leu Trp Ala Asp Pro

500										505										510										
Asp	Arg	Arg	Ala	Ile	Leu	Gly	Val	Asp	Lys	Arg	Thr	Gly	Val	Glu	Pro															
			515									520																		
Ile	Val	Leu	Ala	Glu	Thr	Arg	His	Pro	Pro	Ala	His	Val	Val	Ser	Glu															
			530				535					540																		
Asp	Arg	Asp	Ile	Phe	Ala	Leu	Thr	Gly	Gln	Pro	Asp	Ser	Arg	Asp	Trp															
						550				555					560															
His	Val	Glu	His	Ile	Arg	Ser	Gly	Ala	Ser	Thr	Val	Val	Ala	Asp	Tyr															
				565					570					575																
Gln	Arg	Gln	Leu	Trp	Asp	Arg	Pro	Asp	Met	Val	Leu	Asn	Arg	Arg	Gly															
			580					585					590																	
Leu	Phe	Phe	Thr	Thr	Asn	Asp	Arg	Ile	Leu	Thr	Leu	Ala	Arg	Ser																
			595				600					605																		

<210> 10

<211> 423

<212> PRT

<213> Sorangium cellulosum

<400> 10

Met	Gly	Ala	Leu	Ile	Ser	Val	Ala	Ala	Pro	Gly	Cys	Ala	Leu	Gly	Gly
1				5					10					15	

Ala	Glu	Glu	Glu	Gly	Gln	Pro	Gly	Gln	Asp	Ala	Gly	Ala	Gly	Ala	Leu
			20					25					30		

Ala	Pro	Ala	Arg	Glu	Val	Met	Ala	Ala	Glu	Val	Ala	Ala	Gly	Gln	Met
		35					40					45			

Pro	Gly	Ala	Val	Trp	Leu	Val	Ala	Arg	Gly	Asp	Asp	Val	His	Val	Asp
	50					55					60				

Ala	Val	Gly	Val	Thr	Glu	Leu	Gly	Gly	Ser	Ala	Pro	Met	Arg	Arg	Asp
	65				70					75					80

Thr	Ile	Phe	Arg	Ile	Ala	Ser	Met	Thr	Lys	Ala	Val	Thr	Ala	Thr	Ala
				85					90					95	

Val	Met	Met	Leu	Val	Glu	Glu	Gly	Lys	Leu	Asp	Leu	Asp	Ser	Pro	Val
			100					105					110		

Asp	Arg	Trp	Leu	Pro	Glu	Leu	Ala	Asn	Arg	Lys	Val	Leu	Ala	Arg	Ile
		115					120					125			

Asp	Gly	Pro	Ile	Asp	Glu	Thr	Val	Pro	Ala	Glu	Arg	Pro	Ile	Thr	Val
	130					135					140				

Arg	Asp	Leu	Met	Thr	Phe	Thr	Met	Gly	Phe	Gly	Ile	Ser	Phe	Asp	Ala
	145				150					155					160

Ser	Ser	Pro	Ile	Gln	Arg	Ala	Ile	Asp	Glu	Leu	Gly	Leu	Val	Asn	Ala
				165					170					175	

Gln	Pro	Val	Pro	Met	Thr	Pro	His	Gly	Pro	Asp	Glu	Trp	Ile	Arg	Arg
			180					185					190		

Leu	Gly	Thr	Leu	Pro	Leu	Met	His	Gln	Pro	Gly	Ala	Gln	Trp	Met	Tyr
			195				200					205			

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Asn Thr Gly Ser Leu Val Gln Gly Val Leu Val Gly Arg Ala Ala Asp
 210 215 220
 Gln Gly Phe Asp Ala Phe Val Arg Glu Arg Ile Leu Ala Pro Leu Gly
 225 230 235 240
 Met Arg Asp Thr Asp Phe His Val Pro Ala Asp Lys Leu Ala Arg Phe
 245 250 255
 Ala Gly Cys Gly Tyr Phe Thr Asp Glu Gln Thr Gly Glu Lys Thr Arg
 260 265 270
 Met Asp Arg Asp Gly Ala Glu Ser Ala Tyr Ala Ser Pro Pro Ala Phe
 275 280 285
 Pro Ser Gly Ala Ala Gly Leu Val Ser Thr Val Asp Asp Tyr Leu Leu
 290 295 300
 Phe Ala Arg Met Leu Met Asn Gly Gly Val His Glu Gly Arg Arg Leu
 305 310 315 320
 Leu Ser Ala Ala Ser Val Arg Glu Met Thr Ala Asp His Leu Thr Pro
 325 330 335
 Ala Gln Lys Ala Ala Ser Ser Phe Phe Pro Gly Phe Phe Glu Thr His
 340 345 350
 Gly Trp Gly Tyr Gly Met Ala Val Val Thr Ala Pro Asp Ala Val Ser
 355 360 365
 Glu Val Pro Gly Arg Tyr Gly Trp Asp Gly Gly Phe Gly Thr Ser Trp
 370 375 380
 Ile Asn Asp Pro Gly Arg Glu Leu Ile Gly Ile Val Met Thr Gln Ser
 385 390 395 400
 Ala Gly Phe Leu Phe Ser Gly Ala Leu Glu Arg Phe Trp Arg Ser Val
 405 410 415
 Tyr Val Ala Thr Glu Ser Ala
 420

<210> 11
 <211> 713
 <212> PRT
 <213> Sorangium cellulosum

<400> 11
 Met His Gly Leu Thr Glu Arg Gln Val Leu Leu Ser Leu Val Thr Leu
 1 5 10 15
 Ala Leu Ile Leu Val Thr Ala Arg Ala Ser Gly Glu Leu Ala Arg Arg
 20 25 30
 Leu Arg Gln Pro Glu Val Leu Gly Glu Leu Phe Gly Gly Val Val Leu
 35 40 45
 Gly Pro Ser Val Val Gly Ala Leu Ala Pro Gly Phe His Arg Ala Leu
 50 55 60
 Phe Gln Glu Pro Ala Val Gly Val Val Leu Ser Gly Ile Ser Trp Ile
 65 70 75 80
 Gly Ala Leu Leu Leu Leu Leu Met Ala Gly Ile Glu Val Asp Val Gly
 85 90 95

Ile Leu Arg Lys Glu Ala Arg Pro Gly Ala Leu Ser Ala Leu Gly Ala
 100 105 110
 Ile Ala Pro Pro Leu Ala Ala Gly Ala Ala Phe Ser Ala Leu Val Leu
 115 120 125
 Asp Arg Pro Leu Pro Ser Gly Leu Phe Leu Gly Ile Val Leu Ser Val
 130 135 140
 Thr Ala Val Ser Val Ile Ala Lys Val Leu Ile Glu Arg Glu Ser Met
 145 150 155 160
 Arg Arg Ser Tyr Ala Gln Val Thr Leu Ala Ala Gly Val Val Ser Glu
 165 170 175
 Val Ala Ala Trp Val Leu Val Ala Met Thr Ser Ser Ser Tyr Gly Ala
 180 185 190
 Ser Pro Ala Leu Ala Val Ala Arg Ser Ala Leu Leu Ala Ser Gly Phe
 195 200 205
 Leu Leu Phe Met Val Leu Val Gly Arg Arg Leu Thr His Leu Ala Met
 210 215 220
 Arg Trp Val Ala Asp Ala Thr Arg Val Ser Lys Gly Gln Val Ser Leu
 225 230 235 240
 Val Leu Val Leu Thr Phe Leu Ala Ala Ala Leu Thr Gln Arg Leu Gly
 245 250 255
 Leu His Pro Leu Leu Gly Ala Phe Ala Leu Gly Val Leu Leu Asn Ser
 260 265 270
 Ala Pro Arg Thr Asn Arg Pro Leu Leu Asp Gly Val Gln Thr Leu Val
 275 280 285
 Ala Gly Leu Phe Ala Pro Val Phe Phe Val Leu Ala Gly Met Arg Val
 290 295 300
 Asp Val Ser Gln Leu Arg Thr Pro Ala Ala Trp Gly Thr Val Ala Leu
 305 310 315 320
 Leu Leu Ala Thr Ala Thr Ala Ala Lys Val Val Pro Ala Ala Leu Gly
 325 330 335
 Ala Arg Leu Gly Gly Leu Arg Gly Ser Glu Ala Ala Leu Val Ala Val
 340 345 350
 Gly Leu Asn Met Lys Gly Gly Thr Asp Leu Ile Val Ala Ile Val Gly
 355 360 365
 Val Glu Leu Gly Leu Leu Ser Asn Glu Ala Tyr Thr Met Tyr Ala Val
 370 375 380
 Val Ala Leu Val Thr Val Thr Ala Ser Pro Ala Leu Leu Ile Trp Leu
 385 390 395 400
 Glu Lys Arg Ala Pro Pro Thr Gln Glu Glu Ser Ala Arg Leu Glu Arg
 405 410 415
 Glu Glu Ala Ala Arg Arg Ala Tyr Ile Pro Gly Val Glu Arg Ile Leu
 420 425 430
 Val Pro Ile Val Ala His Ala Leu Pro Gly Phe Ala Thr Asp Ile Val
 435 440 445

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Glu Ser Ile Val Ala Ser Lys Arg Lys Leu Gly Glu Thr Val Asp Ile
 450 455 460
 Thr Glu Leu Ser Val Glu Gln Gln Ala Pro Gly Pro Ser Arg Ala Ala
 465 470 475 480
 Gly Glu Ala Ser Arg Gly Leu Ala Arg Leu Gly Ala Arg Leu Arg Val
 485 490 495
 Gly Ile Trp Arg Gln Arg Arg Glu Leu Arg Gly Ser Ile Gln Ala Ile
 500 505 510
 Leu Arg Ala Ser Arg Asp His Asp Leu Leu Val Ile Gly Ala Arg Ser
 515 520 525
 Pro Ala Arg Ala Arg Gly Met Ser Phe Gly Arg Leu Gln Asp Ala Ile
 530 535 540
 Val Gln Arg Ala Glu Ser Asn Val Leu Val Val Val Gly Asp Pro Pro
 545 550 555 560
 Ala Ala Glu Arg Ala Ser Ala Arg Arg Ile Leu Val Pro Ile Ile Gly
 565 570 575
 Leu Glu Tyr Ser Phe Ala Ala Ala Asp Leu Ala Ala His Val Ala Leu
 580 585 590
 Ala Trp Asp Ala Glu Leu Val Leu Leu Ser Ser Ala Gln Thr Asp Pro
 595 600 605
 Gly Ala Val Val Trp Arg Asp Arg Glu Pro Ser Arg Val Arg Ala Val
 610 615 620
 Ala Arg Ser Val Val Asp Glu Ala Val Phe Arg Gly Arg Arg Leu Gly
 625 630 635 640
 Val Arg Val Ser Ser Arg Val His Val Gly Ala His Pro Ser Asp Glu
 645 650 655
 Ile Thr Arg Glu Leu Ala Arg Ala Pro Tyr Asp Leu Leu Val Leu Gly
 660 665 670
 Cys Tyr Asp His Gly Pro Leu Gly Arg Leu Tyr Leu Gly Ser Thr Val
 675 680 685
 Glu Ser Val Val Val Arg Ser Arg Val Pro Val Ala Leu Leu Val Ala
 690 695 700
 His Gly Gly Thr Arg Glu Gln Val Arg
 705 710

<210> 12
 <211> 126
 <212> PRT
 <213> Sorangium cellulosum

<400> 12
 Met Asp Lys Pro Ile Gly Arg Thr Arg Cys Ala Ile Ala Glu Gly Tyr
 1 5 10 15
 Ile Pro Gly Gly Ser Asn Gly Pro Glu Pro Gln Met Thr Ser His Glu
 20 25 30
 Thr Ala Cys Leu Leu Asn Ala Ser Asp Arg Asp Ala Gln Val Ala Ile

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35 40 45
 Thr Val Tyr Phe Ser Asp Arg Asp Pro Ala Gly Pro Tyr Arg Val Thr
 50 55 60
 Val Pro Ala Arg Arg Thr Arg His Val Arg Phe Asn Asp Leu Thr Glu
 65 70 75 80
 Pro Glu Pro Ile Pro Arg Asp Thr Asp Tyr Ala Ser Val Ile Glu Ser
 85 90 95
 Asp Ala Pro Ile Val Val Gln His Thr Arg Leu Asp Ser Arg Gln Ala
 100 105 110
 Glu Asn Ala Leu Leu Ser Thr Ile Ala Tyr Thr Asp Arg Glu
 115 120 125

<210> 13
 <211> 149
 <212> PRT
 <213> Sorangium cellulosum

<400> 13
 Met Lys His Val Asp Thr Gly Arg Arg Phe Gly Arg Arg Ile Gly His
 1 5 10 15
 Thr Leu Gly Leu Leu Ala Ser Met Ala Leu Ala Gly Cys Gly Gly Pro
 20 25 30
 Ser Glu Lys Thr Val Gln Gly Thr Arg Leu Ala Pro Gly Ala Asp Ala
 35 40 45
 Arg Val Thr Ala Asp Val Asp Pro Asp Ala Ala Thr Thr Arg Leu Ala
 50 55 60
 Val Asp Val Val His Leu Ser Pro Pro Glu Arg Leu Glu Ala Gly Ser
 65 70 75 80
 Glu Arg Phe Val Val Trp Gln Arg Pro Ser Pro Glu Ser Pro Trp Arg
 85 90 95
 Arg Val Gly Val Leu Asp Tyr Asn Ala Asp Ser Arg Arg Gly Lys Leu
 100 105 110
 Ala Glu Thr Thr Val Pro Tyr Ala Asn Phe Glu Leu Leu Ile Thr Ala
 115 120 125
 Glu Lys Gln Ser Ser Pro Gln Ser Pro Ser Ser Ala Ala Val Ile Gly
 130 135 140
 Pro Thr Ser Val Gly
 145

<210> 14
 <211> 184
 <212> PRT
 <213> Sorangium cellulosum

<400> 14
 Val Thr Ser Glu Glu Val Pro Gly Ala Ala Leu Gly Ala Gln Ser Ser
 1 5 10 15
 Leu Val Arg Ala Gln His Ala Ala Arg His Val Arg Pro Cys Thr Arg
 20 25 30

Ala Glu Glu Pro Pro Ala Leu Met His Gly Leu Thr Glu Arg Gln Val
 35 40 45

Leu Leu Ser Leu Val Ala Leu Ala Leu Val Leu Leu Thr Ala Arg Ala
 50 55 60

Phe Gly Glu Leu Ala Arg Arg Leu Arg Gln Pro Glu Val Leu Gly Glu
 65 70 75 80

Leu Phe Gly Gly Val Val Leu Gly Pro Ser Val Val Gly Ala Leu Ala
 85 90 95

Pro Gly Phe His Arg Val Leu Phe Gln Asp Pro Ala Val Gly Val Val
 100 105 110

Leu Ser Gly Ile Ser Trp Ile Gly Ala Leu Val Leu Leu Leu Met Ala
 115 120 125

Gly Ile Glu Val Asp Val Ser Ile Leu Arg Lys Glu Ala Arg Pro Gly
 130 135 140

Ala Leu Ser Ala Leu Gly Ala Ile Ala Pro Pro Leu Arg Thr Pro Gly
 145 150 155 160

Pro Leu Val Gln Arg Met Gln Gly Ala Phe Thr Trp Asp Leu Asp Val
 165 170 175

Ser Pro Arg Arg Ser Ala Gln Ala
 180

<210> 15
 <211> 145
 <212> PRT
 <213> Sorangium cellulosum

<400> 15
 Val Asn Ala Pro Cys Met Arg Cys Thr Ser Gly Pro Gly Val Arg Ser
 1 5 10 15

Gly Gly Ala Ile Ala Pro Ser Ala Glu Ser Ala Pro Gly Arg Ala Ser
 20 25 30

Leu Arg Arg Met Leu Thr Ser Thr Ser Ile Pro Ala Met Ser Ser Arg
 35 40 45

Thr Ser Ala Pro Ile Gln Glu Met Pro Glu Ser Thr Thr Pro Thr Ala
 50 55 60

Gly Ser Trp Lys Arg Thr Arg Trp Asn Pro Gly Ala Ser Ala Pro Thr
 65 70 75 80

Thr Asp Gly Pro Ser Thr Thr Pro Pro Lys Ser Ser Pro Ser Thr Ser
 85 90 95

Gly Trp Arg Ser Arg Arg Ala Ser Ser Pro Lys Ala Arg Ala Val Arg
 100 105 110

Arg Thr Ser Ala Arg Ala Thr Ser Glu Ser Arg Thr Cys Arg Ser Val
 115 120 125

Arg Pro Cys Ile Arg Ala Gly Gly Ser Ser Ala Arg Val Gln Gly Arg
 130 135 140

Thr

145

<210> 16
 <211> 185
 <212> PRT
 <213> Sorangium cellulosum

<400> 16
 Val Leu Ala Pro Pro Ala Asp Ile Arg Pro Pro Ala Ala Ala Gln Leu
 1 5 10 15
 Glu Pro Asp Ser Pro Asp Asp Glu Ala Asp Glu Ala Asp Glu Ala Leu
 20 25 30
 Arg Pro Phe Arg Asp Ala Ile Ala Ala Tyr Ser Glu Ala Val Arg Trp
 35 40 45
 Ala Glu Ala Ala Gln Arg Pro Arg Leu Glu Ser Leu Val Arg Leu Ala
 50 55 60
 Ile Val Arg Leu Gly Lys Ala Leu Asp Lys Val Pro Phe Ala His Thr
 65 70 75 80
 Thr Ala Gly Val Ser Gln Ile Ala Gly Arg Leu Gln Asn Asp Ala Val
 85 90 95
 Trp Phe Asp Val Ala Ala Arg Tyr Ala Ser Phe Arg Ala Ala Thr Glu
 100 105 110
 His Ala Leu Arg Asp Ala Ala Ser Ala Met Glu Ala Leu Ala Ala Gly
 115 120 125
 Pro Tyr Arg Gly Ser Ser Arg Val Ser Ala Ala Val Gly Glu Phe Arg
 130 135 140
 Gly Glu Ala Ala Arg Leu His Pro Ala Asp Arg Val Pro Ala Ser Asp
 145 150 155 160
 Gln Gln Ile Leu Thr Ala Leu Arg Ala Ala Glu Arg Ala Leu Ile Ala
 165 170 175
 Leu Tyr Thr Ala Phe Ala Arg Glu Glu
 180 185

<210> 17
 <211> 146
 <212> PRT
 <213> Sorangium cellulosum

<400> 17
 Met Ala Asp Ala Ala Ser Arg Ser Ala Cys Ser Val Ala Ala Arg Lys
 1 5 10 15
 Leu Ala Tyr Arg Ala Ala Thr Ser Asn Gln Thr Ala Ser Phe Trp Ser
 20 25 30
 Leu Pro Ala Ile Trp Glu Thr Pro Ala Val Val Cys Ala Lys Gly Thr
 35 40 45
 Leu Ser Ser Ala Leu Pro Ser Arg Thr Ile Ala Ser Arg Thr Arg Leu
 50 55 60
 Ser Ser Arg Gly Arg Cys Ala Ala Ser Ala His Arg Thr Ala Ser Glu
 65 70 75 80

[illegible]

<210>	18
<211>	288
<212>	PRT
<213>	Sorangium cellulosum

<400>	18
Val Thr Val Ser Ser Met Pro Arg Ser Trp Ser Ser Arg Val Arg Thr	1 5 10 15
Val Val Thr Ala Leu Gly Cys Ala Arg Arg Leu Ser Gly Ser Ile Ser	20 25 30
Arg Leu Arg Arg His Pro Glu Ala Gly Arg Ala Pro Arg Ser Arg Leu	35 40 45
Arg Ala Trp Arg Arg Leu Pro Gln His Ile Ser Ser Pro Trp Arg His	50 55 60
Leu Pro Pro Gly Ala Arg Val Gly Thr Ser Cys Pro Ala Asp Arg Arg	65 70 75 80
Ile Leu Pro Ser His Arg Thr Ala Asp Leu Gly Thr Ser Gly Gly Thr	85 90 95
Leu Val Ala Arg Met Ser Gly His Val Ala Arg Asn Pro His Ala Ala	100 105 110
Val Leu Val Gly Asp Gly Ser Ala Arg Gly Arg Arg Arg Leu Ser Asn	115 120 125
Arg Arg Ala Glu Arg Arg Val Ser Asp Val Thr Cys Arg Glu Gly Gly	130 135 140
Glu Ala Met Gln Lys Ile Ala Gly Lys Leu Val Val Gly Leu Ile Ser	145 150 155 160
Val Ser Gly Met Ser Leu Leu Ala Ala Cys Gly Gly Glu Lys Arg Ser	165 170 175
Gly Gly Glu Ala Gln Thr Pro Gly Gly Ala Gln Gly Glu Ala Pro Val	180 185 190
Pro Val Gly Ser Ala Val Asp Ser Ile Val Ala Ala Arg Cys Asp Arg	195 200 205
Glu Ala Arg Cys Asn Asn Ile Gly Gln Asp Arg Glu Tyr Ser Ser Lys	210 215 220
Asp Ala Cys Ser Asn Lys Ile Arg Ser Glu Trp Arg Asp Glu Leu Thr	

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225	230	235	240
Phe Gly Glu Cys	Pro Gly Gly Ile Asp	Ala Lys Gln Leu Asn	Glu Cys
	245	250	255
Leu Glu Gly Ile	Arg Asn Glu Gly Cys	Gly Asn Pro Phe Asp	Thr Leu
	260	265	270
Gly Arg Val Val	Ala Cys Arg Ser Ser	Asp Leu Cys Arg	Asp Ala Arg
	275	280	285

<210> 19
 <211> 288
 <212> PRT
 <213> Sorangium cellulosum

<400> 19
 Val Thr Val Ser Ser Met Pro Arg Ser Trp Ser Ser Arg Val Arg Thr
 1 5 10 15
 Val Val Thr Ala Leu Gly Cys Ala Arg Arg Leu Ser Gly Ser Ile Ser
 20 25 30
 Arg Leu Arg Arg His Pro Glu Ala Gly Arg Ala Pro Arg Ser Arg Leu
 35 40 45
 Arg Ala Trp Arg Arg Leu Pro Gln His Ile Ser Ser Pro Trp Arg His
 50 55 60
 Leu Pro Pro Gly Ala Arg Val Gly Thr Ser Cys Pro Ala Asp Arg Arg
 65 70 75 80
 Ile Leu Pro Ser His Arg Thr Ala Asp Leu Gly Thr Ser Gly Gly Thr
 85 90 95
 Leu Val Ala Arg Met Ser Gly His Val Ala Arg Asn Pro His Ala Ala
 100 105 110
 Val Leu Val Gly Asp Gly Ser Ala Arg Gly Arg Arg Arg Leu Ser Asn
 115 120 125
 Arg Arg Ala Glu Arg Arg Val Ser Asp Val Thr Cys Arg Glu Gly Gly
 130 135 140
 Glu Ala Met Gln Lys Ile Ala Gly Lys Leu Val Val Gly Leu Ile Ser
 145 150 155 160
 Val Ser Gly Met Ser Leu Leu Ala Ala Cys Gly Gly Glu Lys Arg Ser
 165 170 175
 Gly Gly Glu Ala Gln Thr Pro Gly Gly Ala Gln Gly Glu Ala Pro Val
 180 185 190
 Pro Val Gly Ser Ala Val Asp Ser Ile Val Ala Ala Arg Cys Asp Arg
 195 200 205
 Glu Ala Arg Cys Asn Asn Ile Gly Gln Asp Arg Glu Tyr Ser Ser Lys
 210 215 220
 Asp Ala Cys Ser Asn Lys Ile Arg Ser Glu Trp Arg Asp Glu Leu Thr
 225 230 235 240

<400> 21
Val Arg Arg Ser Arg Trp Gln Met Lys His Val Asp Thr Gly Arg Arg
1 5 10 15
Val Gly Arg Arg Ile Gly Leu Thr Leu Gly Leu Leu Ala Ser Met Ala
20 25 30
Leu Ala Gly Cys Gly Gly Pro Ser Glu Lys Ile Val Gln Gly Thr Arg
35 40 45
Leu Ala Pro Gly Ala Asp Ala His Val Ala Ala Asp Val Asp Pro Asp

50

55

60

Ala Ala Thr Thr Arg Leu Ala Val Asp Val Val His Leu Ser Pro Pro
 65 70 75 80
 Glu Arg Ile Glu Ala Gly Ser Glu Arg Phe Val Val Trp Gln Arg Pro
 85 90 95
 Ser Ser Glu Ser Pro Trp Gln Arg Val Gly Val Leu Asp Tyr Asn Ala
 100 105 110
 Ala Ser Arg Arg Gly Lys Leu Ala Glu Thr Thr Val Pro His Ala Asn
 115 120 125
 Phe Glu Leu Leu Ile Thr Val Glu Lys Gln Ser Ser Pro Gln Ser Pro
 130 135 140
 Ser Ser Ala Ala Val Ile Gly Pro Thr Ser Val Gly
 145 150 155

<210> 22
 <211> 305
 <212> PRT
 <213> Sorangium cellulosum

<400> 22
 Met Glu Lys Glu Ser Arg Ile Ala Ile Tyr Gly Ala Ile Ala Ala Asn
 1 5 10 15
 Val Ala Ile Ala Ala Val Lys Phe Ile Ala Ala Ala Val Thr Gly Ser
 20 25 30
 Ser Ala Met Leu Ser Glu Gly Val His Ser Leu Val Asp Thr Ala Asp
 35 40 45
 Gly Leu Leu Leu Leu Leu Gly Lys His Arg Ser Ala Arg Pro Pro Asp
 50 55 60
 Ala Glu His Pro Phe Gly His Gly Lys Glu Leu Tyr Phe Trp Thr Leu
 65 70 75 80
 Ile Val Ala Ile Met Ile Phe Ala Ala Gly Gly Gly Val Ser Ile Tyr
 85 90 95
 Glu Gly Ile Leu His Leu Leu His Pro Arg Gln Ile Glu Asp Pro Thr
 100 105 110
 Trp Asn Tyr Val Val Leu Gly Ala Ala Ala Val Phe Glu Gly Thr Ser
 115 120 125
 Leu Ile Ile Ser Ile His Glu Phe Lys Lys Lys Asp Gly Gln Gly Tyr
 130 135 140
 Leu Ala Ala Met Arg Ser Ser Lys Asp Pro Thr Thr Phe Thr Ile Val
 145 150 155 160
 Leu Glu Asp Ser Ala Ala Leu Ala Gly Leu Thr Ile Ala Phe Leu Gly
 165 170 175
 Val Trp Leu Gly His Arg Leu Gly Asn Pro Tyr Leu Asp Gly Ala Ala
 180 185 190
 Ser Ile Gly Ile Gly Leu Val Leu Ala Ala Val Ala Val Phe Leu Ala
 195 200 205

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Ser Gln Ser Arg Gly Leu Leu Val Gly Glu Ser Ala Asp Arg Glu Leu
 210 215 220
 Leu Ala Ala Ile Arg Ala Leu Ala Ser Ala Asp Pro Gly Val Ser Ala
 225 230 235 240
 Val Gly Arg Pro Leu Thr Met His Phe Gly Pro His Glu Val Leu Val
 245 250 255
 Val Leu Arg Ile Glu Phe Asp Ala Ala Leu Thr Ala Ser Gly Val Ala
 260 265 270
 Glu Ala Ile Glu Arg Ile Glu Thr Arg Ile Arg Ser Glu Arg Pro Asp
 275 280 285
 Val Lys His Ile Tyr Val Glu Ala Arg Ser Leu His Gln Arg Ala Arg
 290 295 300

Ala
 305

<210> 23
 <211> 135
 <212> PRT
 <213> Sorangium cellulosum

<400> 23
 Val Gln Thr Ser Ser Phe Asp Ala Arg Tyr Ala Gly Cys Lys Ser Ser
 1 5 10 15
 Arg Arg Ile Ala Arg Ser Gly Ser Ala Gly Ala Arg Ala Gly Arg Ala
 20 25 30
 His Glu Gly Ala Ala Ser Ala Gly Phe Glu Gly Gly Asp Val Met Arg
 35 40 45
 Lys Ala Arg Ala His Gly Ala Met Leu Gly Gly Arg Asp Asp Gly Trp
 50 55 60
 Arg Arg Gly Leu Pro Gly Ala Gly Ala Leu Arg Ala Ala Leu Gln Arg
 65 70 75 80
 Gly Arg Ser Arg Asp Leu Ala Arg Arg Arg Leu Ile Ala Ser Val Ser
 85 90 95
 Leu Ala Gly Gly Ala Ser Met Ala Val Val Ser Leu Phe Gln Leu Gly
 100 105 110
 Ile Ile Glu Arg Leu Pro Asp Pro Pro Leu Pro Gly Phe Asp Ser Ala
 115 120 125
 Lys Val Thr Ser Ser Asp Ile
 130 135

<210> 24
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: universal
 reverse primer

<400> 24

ggaaacagct atgaccatg

19

<210> 25

<211> 17

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: universal
forward primer

<400> 25

gtaaaacgac ggccagt

17

<210> 26

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
NH24 end "B"

<400> 26

gtgactggcg cctggaatct gcatgagc

28

<210> 27

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer NH2
end "A"

<400> 27

agcgggagct tgctagacat tctgtttc

28

<210> 28

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer NH2
end "B"

<400> 28

gacgcgcctc gggcagcgcc ccaa

24

<210> 29

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
pEPO15-NH6 end "B"

<400> 29

caccgaagcg tcgatctggt ccac

25

<210> 30

<211> 25

<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
pEPO15H2.7 end "A"

<400> 30
cggtcagatc gacgacgggc ttcc

25